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(54) Title: CRYSTAL STRUCTURE OF A DEACETYLASE AND INHIBITORS THEREOF

(57) Abstract: The present invention provides three-dimensional structural information from the hyperthermophilic bacterium Aquifex aeolicus which is a histone deacetylase-like protein (HDLP). HDLP shares 35.2% amino acid sequence identity with human histone deacetylase (HDAC1). The present invention further provides three-dimensional structural information of HDLP bound by inhibitor molecules. The three-dimensional structural information of the present invention is useful to design, isolate and screen deacetylase inhibitor compounds capable of inhibiting HDLP, HDAC family members and HDLP-related molecules. The invention also relates to nucleic acids encoding a mutant HDLP which facilitates the determination of the three-dimensional structure of HDLP in the presence of a zinc atom.



Crystal Structure of a Deacetylase and Inhibitors Thereof

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This application claims priority of U.S. Provisional Application No. 60/152,753, filed September 8, 1999, the contents of which are hereby incorporated by reference.

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This invention has been made with government support under National Institutes of Health Grant No. RO1 CA-65698. Accordingly, the U.S. Government may have certain rights in the invention.

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Throughout this application, various publications are referenced by author, date and citation. The disclosures of these publications in their entireties are hereby incorporated by reference into this application in order to more fully describe the state of the art as known to those skilled therein as of the date of the invention described and claimed herein.

Introduction

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The present invention relates to a histone deacetylase homologue from the hyperthermophilic bacterium Aquifex aeolicus, HDLP (histone deacetylase like protein; also known as AcuCl), which shares 35.2 % sequence identity with human histone deacetylase (HDACl), that can be co-crystallized with an inhibitory ligand, and more particularly, to the detailed crystallographic data obtained from said co-crystallization which is disclosed herein. The invention also relates to methods of using the crystal structure and x-ray crystallographic coordinates of the apo-HDLP and

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inhibitor-bound HDLP to design, isolate and screen compounds which bind to and inhibit the active site of HDLP and HDLP-related proteins, such as those proteins belonging to the HDAC family, including HDAC1.

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Background of the Invention

The reversible modification of histones by acetylation is associated with changes in nucleosome conformation and chromatin structure, and plays an important role in the regulation of gene expression (reviewed in Davie and Chadee, 1998, J. Cell Biochem. Suppl. 30-31:203-213). The histone acetylase and deacetylase enzymes that carry out these modifications are involved in many cellular processes such as cell cycle progression and differentiation, and their deregulation is associated with several types of human cancer (reviewed in Kouzarides, 1999, Curr. Opin. Genet. Dev. 2:40-48; Hassig et al., 1997, Chem. Biol. 4:783-789; Fenrick and Heibert, 1998, J. Cell. Biochem. Suppl. 30-31:194-202).

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Recently, several experimental antitumor compounds, such as trichostatin A (TSA), trapoxin, suberoylanilide hydroxamic acid (SAHA), and phenylbutyrate have been shown to act, at least in part, by inhibiting histone deacetylases. Richon et al., 1998, Proc. Natl. Acad. Sci., USA 95:3003-3007; Yoshida et al., 1990, J. Biol. Chem. 265:17174-17179; Kijima et al., 1993, J. Biol. Chem. 268:22429-22435. Additionally, diallyl sulfide and related molecules (Lea et al., 1999, Int. J. Oncol. 2:347-352), oxamflatin (Kim et al., 1999, Oncogene 15:2461-2470), MS-27-275, a synthetic benzamide derivative (Saito et al., 1999, Proc. Natl. Acad. Sci. 96:4592-4597),

butarate derivatives (Lea and Tulsyan, 1995, Anticancer Res. 15:879-883), FR901228 (Nokajima et al., 1998, Exp. Cell Res. 241:126-133), depudecin (Kwon et al., 1998, Proc. Natl. Acad. Sci. USA 95:3356-3361) and m-carboxysinnamic acid bishydroxamide (CBHA; Richon et al., Proc. Natl. Acad. Sci. USA 95:3003-3007) have been shown to inhibit histone deacetylases. In vitro, these compounds can inhibit the growth of fibroblast cells by causing cell cycle arrest in the G1 and G2 phases (Richon et al., 1996, Proc. Natl. Acad. Sci. USA 93:5705-5708; Kim et al., 1999, Oncogene 18:2461-2470; Yoshida et al., 1995, Bioessays 17:423-430; Yoshida & Beppu, 1988, Exp. Cell. Res. 177:122-131), and can the terminal differentiation and loss of to transforming potential of a variety of transformed cell lines. Richon et al., 1996, Proc. Natl. Acad. Sci. USA 93:5705-5708; Kim et al., 1999, Oncogene 18:2461-2470; Yoshida et al., 1987, Cancer Res. 47:3688-3691. In vivo, phenylbutyrate is effective in the treatment of acute promyelocytic leukemia in conjunction with retinoic acid. Warrell et al., 1998, J. Natl. Cancer Inst. 90:1621-1625. SAHA is effective in preventing the formation of mammary tumors in rats, and lung tumors in mice. Desai et al., 1999, Proc. AACR 40: abstract #2396; Cohen et al., Cancer Res., submitted.

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Histone deacetylases catalyze the removal of acetyl groups from the e-amino groups of lysine residues clustered near the N-terminus of nucleosomal histones, and this process is associated with transcriptional repression (reviewed in Struhl, 1998, Genes Dev. 12:599-606). Deletion of the yeast histone deacetylase gene, rpd3, or its pharmacological

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inactivation with trichostatin A reduces the transcriptional repression in a subset of promoters, such as those of Ume6-regulated genes. Kadosh & Struhl, 1998, Mol. Cell. Biol. 18:5121-5127. This is accompanied by the increased acetylation of H4 histones in the repressed promoter and its vicinity, but has no effect on histones at promoter distal regions. Kadosh & Struhl, 1998, Mol. Cell. Biol. 18:5121-5127; Rundlett et al., 1998, Nature 392:831-835.

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Histone deacetylases are recruited to specific promoters by associating with DNA-binding transcriptional repressors, either directly or through co-repressors that bridge the deacetylase to the transcriptional repressors. For example, the Mad and Ume6 repressors bind to the co-repressor Sin3A (Laherty et al., 1997, Cell 89:349-356; Hassig et al., 1997, Cell 89:341-347; Kadosh & Struhl, 1997, Cell 89:365-371), and the nuclear receptors bind N-CoR and the related SMRT co-repressors. Nagy et al., 1997, Cell 89:373-380; Alland et al, 1997, Nature 387:49-55; Heinzel et al, 1997, Nature 387:43-48.

The deregulation of histone deacetylase recruitment appears to be one of the mechanisms through which these enzymes contribute to tumorigenesis. In acute promyelocytic leukemia (APL), chromosomal translocations fuse the retinoic acid receptor-α (RARα) to either PLZF or to PML. These fusion oncoproteins have aberrant transcriptional repression activity resulting, in part, through the recruitment of a co-repressor and, in turn, HDACs. Grignani et al, 1998, Nature 391:815-818; Lin et al., 1998, Nature 391:811-814. Treatment of PLZF-RARα APL cells with TSA enhances their

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responsiveness to retinoic acid-induced differentiation. Grignani et al, 1998, Nature 391:815-818; Lin et al., 1998, Nature 391:811-814.

histone deacetylases comprise a large family of 5 proteins, conserved from yeast to man, and are divided into two related classes. Class I is characterized by human HDAC1, 2, 3 (Taunton et al., 1996, Science 272:408-411; Yang et al., 1996, Proc. Natl. Acad. Sci. USA 93:12845-12850; Emiliani et al., 1998, Proc. Natl. Acad. Sci. USA 95:2795-10 2800), and yeast RPD3 (Videl & Gaber, 1991, Mol. Cell. Biol. 11:6317-6327), and class II by the human HDAC4, 5, 6 (Grozinger et al., 1999, Proc. Natl. Acad. Sci. USA 96:4868-4873; Fischle, et al., 1999, J. Biol. Chem. <u>274</u>:11713-11720), and yeast HDA1 (Rundlett et al., 1996, Proc. Natl. 15 Acad. Sci. USA <u>93</u>:14503-14508). The two classes share a ~390 amino acid region of sequence similarity, comprising the deacetylase core, but are divergent outside this region. The histone deacetylase genes belong to an even larger superfamily (Leipe & Landsman, 1997, Nucleic Acids Res. 20 that contains the prokaryotic 25:3693-3697) utilization proteins (AcuC; 28.1% sequence identity to HDAC1), and the prokaryotic acetylpolyamine amidohydrolases (APAH; 15.0 % sequence identity to HDAC1). The enzymatic activity of AcuC is not clear, but its disruption reduces 25 the ability of B. subtilis to breakdown acetoin and utilize it as a carbon source. Grundy et al., 1993, Mol. Microbiol. 10:259-271. APAHs catalyze the deacetylation of polyamines by cleaving a non-peptide amide bond (reviewed in Leipe & Landsman, 1997, Nucleic Acids Res. 25:3693-3697). 30

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It is useful to address the questions of how HDACs and HDACrelated proteins catalyze the deacetylation of histones and how the above-referenced compounds, particularly those compounds with antitumor activity, inhibit this activity in order to better understand the mechanism of inhibition of HDACs and to facilitate discovery of additional useful compounds which may inhibit this activity. To this end, the present invention has determined the three dimensional structure of a HDAC1-like protein from the thermophilic bacterium Aquifex aeolicus, herein after HDLP. determination of the nucleic acid coding sequence of HDLP was described by Deckert et al., 1998, Nature 392:353-358. encoded 375 residue protein, whose sequence was determined from the nucleic acid encoding sequence, shares 35.2% amino acid sequence identity with HDAC1, deacetylates histones in vitro, and is inhibited by TSA, SAHA and several other HDAC inhibitors. The determination of the threedimensional structure of HDLP is useful in the design, identification and screening of new HDAC family inhibitory compounds which are useful for the inhibition of cell growth both in vivo and in vitro.

Summary of the Invention

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In general, it is the object of the present invention to provide detailed three-dimensional structural information for a family of proteins known as histone deacetylases (HDAC), and particularly a homologue from the hyperthermophilic bacterium Aquifex aeolicus HDLP (histone deacetylase-like protein) which shares 35.2 % sequence identity with human histone deacetylase (HDAC1). It is also an object of the present invention to provide three-

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dimensional structural information of an HDLP bound to an inhibitory compound.

In one embodiment of the invention, three-dimensional structure information is obtained from a crystal of wildtype HDLP (SEQ ID NO:1) (the nucleic acid encoding wild-type In a further embodiment of the HDLP is SEQ ID NO:2). invention, three-dimensional information is obtained from a mutant HDLP comprising two mutations (1) cysteine 75 to a serine and (2) cysteine 77 to a serine (Cys75Ser/Cys77Ser double mutant; SEQ ID NO:3) (the nucleic acid encoding HDLP Cys75Ser/Cys77Ser double mutant is SEQ ID NO:4). The HDLP invention facilitates of the present mutant determination of three-dimensional structural information of HDLP bound to a zinc atom at its zinc atom-binding site.

In a preferred embodiment of the invention, the threedimensional structural information is obtained from a cocrystal of a protein-inhibitor compound complex that comprises HDLP or HDLP Cys75Ser/Cys77Ser double mutant and trichostatin A (TSA). In another preferred embodiment of the invention the three-dimensional structural information is obtained from a co-crystal of a protein-inhibitor or HDLP HDLP complex that comprises compound Cys75Ser/Cys77Ser suberoylanilide double mutant and hydroxamic acid (SAHA). Any HDLP or HDLP-related protein (e.g. HDAC) inhibitor compound that may be co-crystallized with HDLP may be used to form a co-crystal of the present invention.

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The protein crystals and protein-inhibitory complex cocrystals of the present invention diffract to a high

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resolution limit of at least equal to or greater than 4 angstrom (Å). In a preferred embodiment, the protein crystals and protein-inhibitory complex co-crystals of the present invention diffract to a high resolution limit of greater than 2.5 Å.

A crystal of the present invention may take a variety of forms, all of which are contemplated by the present invention. In a preferred embodiment, the crystal has a space group of C2 with one molecule in the asymmetric unit and with unit dimensions of a = 51.4 Å, b = 93.8 Å, c =78.7 Å and $\beta = 96.9^{\circ}$ (see, e.g., Example 2, below). another preferred embodiment, the crystal has a space group of $P2_12_12_1$ with two molecules in the asymmetric unit and with unit dimensions of a = 53.4 Å, b = 94.4 Å, c = 156.3 Å (see, e.g., Example 2, below). The HDLP structure comprises a parallel β sheet with α helices packing against both faces. At one end of the β sheet, the HDLP has a narrow, tube-like pocket formed by several well-ordered loops. The walls of the pocket are lined with hydrophobic residues and there is a zinc binding site and several polar side chains at the bottom of the pocket. The inhibitory compounds of the present invention bind in the pocket.

The three-dimensional structural information obtained from crystals of HDLP, HDLP Cys75Ser/Cys77Ser double mutant, HDLP Cys75Ser/Cys77Ser double mutant comprising a zinc atom, HDLP comprising an inhibitory compound such as TSA or SAHA, and HDLP Cys75Ser/Cys77Ser double mutant comprising an inhibitor compound such as TSA or SAHA may be employed to solve the structure of any HDLP-related protein (e.g. HDAC) crystal,

or any mutant HDLP-related protein and particularly any wild type or mutant of HDLP-related protein complexed with a ligand, including a substrate or inhibitor compound. If the crystals are in a different space group than the known structure, molecular replacement may be employed to solve the structure, or if the crystals are in the same space group, refinement and difference fourier methods may be employed. The structure of HDLP-related proteins (e.g. HDAC1) comprise no greater than a 2.0 Å root mean square deviation (rmsd) in the positions of the C α atoms for at least 50% or more of the amino acids of the full-length HDLP structure.

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The present invention also provides a nucleic acid molecule encoding an HDLP Cys75Ser/Cys77Ser double mutant having the amino acid sequence of SEQ ID NO:3 and the nucleic acid sequence of SEQ ID NO:4. It is also contemplated by the invention that mutations be made in HDLP-related proteins at cysteine residues, as with the Cys75Ser/Cys77Ser double mutant, in order to facilitate the determination of the structure of said proteins bound to a zinc atom. Additionally, the present invention provides expression vectors which comprise the nucleic acid molecule encoding an HDLP Cys75Ser/Cys77Ser double mutant encoded by the sequence represented by SEQ ID NO:4 operatively linked to expression control sequences.

It is another object of the present invention to provide methods for the design, identification and screening of potential inhibitor compounds of the HDLP/HDAC family. In a preferred embodiment the method for the rational design,

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identification and screening of potential inhibitor compounds for HDLP and HDLP-related proteins (e.g. HDACs) comprising deacetylase activity comprises the steps of: (a) using a three-dimensional structure of an HDLP as defined by the atomic coordinates of the present invention; (b) employing said three-dimensional structure to design or select said potential inhibitor compound; (c) synthesizing and/or selecting said potential inhibitor; (d) contacting said potential inhibitor compound with said enzyme in the presence of acetylated substrate; and (e) determining the percent inhibition of deacetylase activity to determine the inhibitory activity of said potential inhibitor compound. In a further preferred embodiment, the binding properties of said rationally designed inhibitory compound may be determined by a method comprising the steps of: (a) forming a complex comprising said inhibitory compound and HDLP or a HDLP-related protein, (b) co-crystallizing said inhibitory compound-HDLP complex; (c) determining said dimensional structure of said co-crystal through molecular replacement or refinement and difference fourier with the molecular coordinates of HDLP as defined by the present invention; and (d) analyzing the three-dimensional structure to determine the binding characteristics of said potential inhibitor compound.

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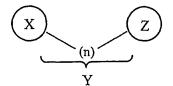
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It is a further object of the present invention to identify a defined class of HDLP/HDAC family inhibitor compounds. The HDLP/HDAC family inhibitor compounds of the present invention are represented by formula (I):

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(I)



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wherein X comprises a cap group which binds to at least one amino acid selected from the group consisting of proline and leucine; Y comprises an aliphatic chain group which binds to at least one amino acid selected from the group consisting of leucine, phenylalanine and glycine; and Z comprises and active site binding group which binds to at least one amino acid selected from the group consisting of aspartic acid, tyrosine and histidine and may further bind to a zinc atom.

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Brief Description of the Drawings

Figure 1 is a table listing the statistics from the X-ray crystallographic analysis of a HDLP crystal, a HDLP-TSA cocrystal, and a HDLP-SAHA co-crystal.

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Figure 2 shows an alignment of various HDAC homologues with percent sequence identity depicted.

Figure 3 shows a graph indicating the histone deacetylase activity of HDLP and HDAC1 and the inhibition of HDLP and HDAC1 by the inhibitors TSA and HC-toxin.

Figure 4 shows (A & B) a schematic representation of the $HDLP-Zn^{2+}-TSA$ complex in two approximately orthogonal views, (C) a topology diagram of HDLP indicating the regions of homology with HDAC1, and (D) a close-up schematic representation of the $HDLP-Zn^{2+}-SAHA$ complex.

Figure 5 shows (A) a schematic representation of a slice through a surface representation of HDLP with the pocket internal cavities and position of the β sheet indicated, (B) a schematic representation of a close-up view of the active site looking down into the pocket in an orientation similar to Figure 4B.

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Figure 6 shows (A) a space-filling representation of TSA in the active site pocket, (B) a closeup stereo view of the structure of the $HDLP-ZN^{2+}-TSA$ complex in a similar orientation to Figure 4B, and (C) a schematic representation of the HDLP-TSA interactions.

Figure 7 shows (A) a schematic representation of the regions of homology shared between HDLP and HDAC1 in an orientation similar to that of Figure 4A, and (B) a detailed schematic representation of the homology shared in the pocket and internal cavity between HDLP and HDAC1 in an orientation similar to that of Figure 4B.

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Figure 8 shows a schematic representation of the proposed catalytic mechanism for the deacetylation of acetylated lysine.

Figure 9 shows a schematic representation of a space filling diagram showing the conserved amino acids in the active site and nearby grooves.

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Figure 10 is the nucleic acid sequence of HDLP from Aquifex aeolicus (SEQ ID NO. 2).

Figure 11 is the amino acid sequence of full length HDLP from Aquifex aeolicus (SEQ ID NO. 1).

Figure 12 is the nucleic acid sequence of the HDLP active site mutant Tyr297Phe (SEQ ID NO. 6).

Figure 13 is the amino acid sequence of the HDLP active site mutant Tyr297Phe (SEQ ID NO. 5).

Figure 14 is the nucleic acid sequence of a double mutant of HDLP from Aquifex aeolicus comprising a Cys75Ser and Cys77Ser mutation (SEQ ID NO. 4).

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Figure 15 is the amino acid sequence of a double mutant of HDLP from Aquifex aeolicus comprising a Cys75Ser and Cys77Ser mutation (SEQ ID NO. 3).

Figure 16-1 to 16-49 lists the atomic structure coordinates for HDLP as derived by X-ray diffraction from a crystal of HDLP.

Figure 17-1 to 17-49 lists the atomic structure coordinates for HDLP Cys75Ser/Cys77Ser double mutant comprising a zinc atom in the active site as derived by X-ray diffraction from a crystal of the HDLP Cys75Ser/Cys77Ser double mutant.

Figure 18-1 to 18-99 lists the atomic structure coordinates for HDLP Cys75Ser/Cys77Ser double mutant as derived by X-ray diffraction from a co-crystal of HDLP complexed with TSA.

Figure 19-1 to 19-48 lists the atomic structure coordinates for HDLP Cys75Ser/Cys77Ser double mutant as derived by X-ray diffraction from a co-crystal of HDLP complexed with SAHA.

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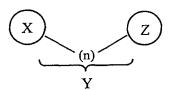
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Detailed Description of the Invention

The present invention provides crystals of a histone deacetylase (HDAC) homologue grown in the presence and absence of a compound capable of inhibiting the histone deacetylase activity of said HDAC homologue. As referred to herein, a HDAC homologue (as well as a HDLP-related protein) is any protein molecule having (a) greater than 15% sequence identity to over the 375 amino acid residues of HDLP; (b) having no more than twenty insertions or deletions for a total of no more than 100 amino acids; and (c) deacetylase activity. Sequence identity is calculated by the program DNAstar[™] using the identity matrix weighing scheme clustal method (DNAstar program, Madison, WI).

A HDLP/HDAC inhibitor compound, as used herein, refers to any compound represented by Formula (I):

(I)



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wherein X comprises a cap group which binds to at least one amino acid selected from the group consisting of tyrosine, proline and leucine; Y comprises an aliphatic chain group from about 5 to about 10 Å, preferably 7Å, which binds to at least one amino acid selected from the group consisting of phenylalanine and glycine; and Z comprises a active site binding group which binds to at least one amino acid selected from the group consisting of aspartic acid, tyrosine and histidine and which may further bind to a zinc atom. The HDAC inhibitory compounds of the present

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invention can inhibit greater than 50% of the histone deacetylase activity of a HDAC homologue or a HDLP-related protein.

To grow the crystals of the present invention, the HDAC and HDAC-inhibitory compound complex are purified to greater than 80% total protein and more preferably purified to greater than 90% total protein. For expression and purification purposes, the full-length HDLP (Genbank accession number AE000719) may be subcloned from Aquifex aeolicus chromosomal DNA preparation by the polymerase chain reaction (PCR) and inserted into an expression vector.

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A large number of vector-host systems known in the art may be used. Possible vectors include, but are not limited to, plasmids or modified viruses, but the vector system must be compatible with the host cell used. Examples of vectors include E. coli bacteriophages such as lambda derivatives, or plasmids such as pBR322 derivatives or pUC plasmid derivatives, e.g., pGEX vectors (Amersham-Pharmacia, Piscataway, New Jersey), pET vectors (Novagen, Madison, WI), pmal-c vectors (Amersham-Pharmacia, Piscataway, New Jersey), pFLAG vectors (Chiang and Roeder, 1993, Pept. Res. 6:62-64), baculovirus vectors (Invitrogen, Carlsbad, CA; Pharmingen, San Diego, CA), etc. The insertion into a cloning vector can, for example, be accomplished by ligating the DNA fragment into a cloning vector which has complementary cohesive termini, by blunt end ligation if no complementary cohesive termini are available or by through nucleotide linkers using techniques standard in the art. Ausubel et al. (eds.), Current Protocols in Molecular

Biology, (1992). Recombinant vectors comprising the nucleic acid of interest may then be introduced into a host cell compatible with the vector (e.g. E. coli, insect cells, mammalian cells, etc.) via transformation, transfection, infection, electroporation, etc. The nucleic acid may also be placed in a shuttle vector which may be cloned and propagated to large quantities in bacteria and then introduced into a eukaryotic cell host for expression. The vector systems of the present invention may provide expression control sequences and may allow for the expression of proteins in vitro.

In a preferred embodiment, the full length HDLP (SEQ ID NO:2) is subcloned from Aquifex aeolicus chromosomal DNA preparation into pGEX4T3 (Amersham-Pharmacia, Piscataway, New Jersey). In order to construct a double mutant comprising a Cys75Ser and Cys77Ser mutation (SEQ ID NO:4), and to construct the HDLP active site mutant Tyr297Phe (SEQ ID NO:5 and SEQ ID NO:6), PCR site directed mutagenesis may be employed with verification by DNA sequencing by methods known to those skilled in the art (see, e.g., Example 1 below). The mutants of the present invention may be subcloned into a suitable expression vector and introduced into a host cell for protein production, as described above.

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The HDLP nucleic acids of the present invention may be subcloned into an expression vector to create an expression construct such that the resultant HDLP molecule which is produced comprises a fusion protein wherein said fusion protein comprises a tag for ease of purification. As referred to herein, a "tag" is any additional amino acids

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which are provided in a protein either c-terminally, nterminally or internally for the ease of purification, for the improvement of production or for any other purpose which may facilitate the goals of the present invention (e.g. to achieve higher levels of production and/or purification). Such tags include tags known to those skilled in the art to be useful in purification such as, but not limited to, his tag, glutathione-s-transferase tag, flag tag, mbp (maltose binding protein) tag, etc. In a preferred embodiment, the wild-type and mutant HDLPs of the present invention are tagged with glutathione-s-transferase (see Example 1 below). In another preferred embodiment, HDAC1 is flag tagged (see Example 1 below). Such tagged proteins may also be engineered to comprise a cleavage site, such as a thrombin, enterokinase or factor X cleavage site, for ease of removal of the tag before, during or after purification. systems which provide a tag and a cleavage site for removal of the tag are particularly useful to make the expression constructs of the present invention.

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The tagged HDLPs and HDACs of the present invention may be purified by immuno-affinity or conventional chromatography, including but not limited to, chromatography employing the (Amersham-Pharmacia, qlutathione-sepharose™ following: Piscataway, New Jersey) or an equivalent resin, nickel or cobalt-purification resins, anion exchange chromatography, cation exchange chromatography, hydrophobic resins, gel antiflag epitope resin, reverse filtration, chromatography, etc. After purification, the HDLP and HDLPinhibitor compound complex may be concentrated to greater than 1 mg/ml for crystallization purposes. In a preferred embodiment HDLP and HDLP-inhibitor complexes

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concentrated to greater than 10 mg/ml for crystallization and in a particularly preferred embodiment, HDLP and HDLP-inhibitor complexes are concentrated to greater than 20 mg/ml.

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In order to determine whether the purified HDLPs of the present invention demonstrate histone deacetylase activity, the purified HDLPs and also any HDLP-related protein may be assayed by any method known to those skilled in the art for the determination of said activity. In a preferred embodiment, the purified HDLPs of the present invention are incubated in the presence of [3H]acetyl-labeled histone substrate (Carmen et al., 1996, J. Biol. Chem. 271:15837-15844) in a buffer suitable for detection of histone deacetylase activity (see Example 3 below); stopping the reaction; extracting the released acetate and measuring said released acetate, as described by Henzel et al. (J. Biol. Chem. 266:21936-21942 (1991); Example 3 below). preferred embodiment, the HDLPs of the present invention are inclubated in the presence of ${\rm ZnCl_2}$ in order to obtain histone deacetylase activity therefrom (Example 3 below).

In another embodiment, the crystals of the present invention comprise purified wild-type HDLP (SEQ ID NO:1) and are grown at room temperature by the hanging-drop vapor-diffusion method from a crystallization solution comprising one or more precipitants selected from the group consisting of isopropanol, polyethylene glycol, and tert butanol (see Example 2 below). The crystallization solution may further comprise one or more salts including salts selected from the group consisting of NaCl and KCl, and one or more buffers

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including buffers selected from the group consisting of Tris (tris(hydroxymethyl)aminomethane and bis-tris propane-Cl (1,3-bis[tris(hydroxymethyl)methyl-amino] propane) (see Example 2 below). The pH of the crystallization solution is preferably between pH 5 to 9, although other pH values are also contemplated by the present invention (see Example 2 below).

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Any crystallization technique known to those skilled in the art may be employed to obtain the crystals of the present invention, including, but not limited to, batch crystallization, vapor diffusion (either by sitting drop or hanging drop) and micro dialysis. Seeding of the crystals in some instances may be required to obtain X-ray quality crystals. Standard micro and/or macro seeding of crystals may therefore be used.

The crystals of the present invention may form in the space group C2 with one molecule in the asymmetric unit and with unit dimensions of a=51.4 Å, b=93.8 Å, c=78.7 Å and $\beta=96.9^{\circ}$ (see Example 2 below). The crystals of the present invention may also form in the space group $P2_12_12_1$ with two molecules in the asymmetric unit and with unit dimensions of a=53.4 Å, b=94.4 Å, c=156.3 Å (see Example 2 below). However, the present invention contemplates crystals which form in any space group including, but not limited to, C2, $P2_1$, $P2_12_12_1$, $P3_121$, $P4_32_12_1$, and $C222_1$. The crystals diffract to a resolution greater than 4 Å, preferably greater than 2.5 Å.

To collect diffraction data from the crystals of the present

invention, the crystals may be flash-frozen in the crystallization buffer employed for the growth of said crystals, however with preferably higher precipitant concentration (see, e.g., Example 2 below). For example, but not by way of limitation, if the precipitant used was 28% PEG 1500, the crystals may be flash frozen in the same crystallization solution employed for said crystal growth wherein the concentration of the precipitant is increased to 35% (see Example 2 below). If the precipitant is not a sufficient cryoprotectant (i.e. a glass is not formed upon glycerol, flash-freezing), cryoprotectants (e.g. molecular weight PEGs, alcohols, etc) may be added to the solution in order to achieve glass formation upon flashfreezing, providing the cryoprotectant is compatible with preserving the integrity of the crystals. The flash-frozen crystals are maintained at a temperature of less than -110°C and preferably less than -150°C during the collection of the crystallographic data by X-ray diffraction. diffraction data may be processed with DENZO and SCALEPACK (Otwinowski & Minor, 1997, Method Ensemble. 276:307-326) but any method known to those skilled in the art may be used to process the X-ray diffraction data.

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In order to determine the atomic structure of HDLP according to the present invention, multiple isomorphous replacement (MIR) analysis, model building and refinement may be performed. For MIR analysis, the crystals may be soaked in heavy-atoms to produce heavy atom derivatives necessary for MIR analysis. As used herein, heavy atom derivative or derivitization refers to the method of producing a chemically modified form of a protein or protein complex

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crystal wherein said protein is specifically bound to a heavy atom within the crystal. In practice a crystal is soaked in a solution containing heavy metal atoms or salts, or organometallic compounds, e.g., lead chloride, gold cyanide, thimerosal, lead acetate, uranyl acetate, mercury chloride, gold chloride, etc, which can diffuse through the crystal and bind specifically to the protein. The location(s) of the bound heavy metal atom(s) or salts can be determined by X-ray diffraction analysis of the soaked This information is used to generate MIR phase crystal. information which is used to construct the three-dimensional structure of the crystallized HDLPs and HDLP-related proteins of the present invention. In a preferred embodiment, the heavy atoms comprise thimerosal, KAu(CN)2 and Pb(Me)3OAc (see Example 2 below). The MIR phases may be calculated by any program known to those skilled in the art and preferably with the program MLPHARE (The CCP4 suite: Programs for computational crystallography, 1994, Crystallogr. D. 50:760-763) and may also use the anomalous diffraction signal from the thimerosal derivative. preferred embodiment, the MIR phases were calculated at 2.5 Å and have a mean figure of merit of 0.55 (see Figure 19 and Example 2 below). The phases may be improved where necessary by solvent flattening by methods known to those skilled in the art including, but not limited to, through the use of the program DM (The CCP4 suite: Programs for computational crystallography, 1994, Acta Crystallogr. D <u>50</u>:760-763).

Thereafter, an initial model of the three-dimensional structure may be built using the program O (Jones et al.,

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1991, Acta Crystallogr. A 47:110-119). The interpretation and building of the structure may be further facilitated by use of the program CNS (Brunger et al., 1998, Acta Crystallogr. D 54:905-921).

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For the determination of the HDLP-inhibitor compound complex structure, if the space group of the HDLP-inhibitor compound complex crystal is different, molecular replacement may be employed using a known structure of apo-HDLP (as referred to herein, apo-HDLP or apo-HDAC is the enzyme which is not complexed with an inhibitor compound) or any known HDLP/inhibitor complex structure whose structure may be determined as described above and below in Example 2. the space group of the HDLP-inhibitor compound crystals is the same, then rigid body refinement and difference fourier may be employed to solve the structure using a known structure of apo-HDLP (as referred to herein, apo-HDLP or apo-HDAC is the enzyme which is not complexed with an inhibitor compound) or any known HDLP/inhibitor complex structure.

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The term "molecular replacement" refers to a method that involves generating a preliminary model of the three-dimensional structure of the HDLP crystals of the present invention whose structure coordinates are unknown prior to the employment of molecular replacement. Molecular replacement is achieved by orienting and positioning a molecule whose structure coordinates are known (in this case the previously determined apo-HDLP) within the unit cell as defined by the X-ray diffraction pattern obtained from an HDLP or HDLP-related protein crystal whose structure is

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unknown so as to best account for the observed diffraction pattern of the unknown crystal. Phases can then be calculated from this model and combined with the observed amplitudes to give an approximate Fourier synthesis of the structure whose coordinates are unknown. This in turn can be subject to any of several forms of refinement to provide a final, accurate structure.

Any method known to the skilled artisan may be employed to determine the structure by molecular replacement. example, the program AMORE (The CCP4 suite: Programs for computational crystallography, 1994, Acta Crystallogr. D. 50:760-763) may be employed to determine the structure of an unknown histone deacetylase +/- an inhibitor by molecular replacement using the apo-HDLP coordinates (Figure 16). For the structure determination of the inhibitory compound TSA, the structure of TSA was obtained from the Cambridge Structural Database (Refcode TRCHST, http://www.ccdc.cam.ac.uk >>) may be employed to define the stereochemical restraints used in the refinement with the program CNS (Brunger et al., 1998, Acta Crystallogr. D <u>54</u>:905-921).

The three-dimensional structural information and the atomic coordinates associated with said structural information of HDLP are useful for solving the structure of crystallized proteins which belong to the HDAC family by molecular replacement. Similarly, any structure of a crystallized protein which is thought to be similar in structure based on function or sequence similarity or identity to HDLP may be solved by molecular replacement with the HDLP structural

information of the present invention. The structure of HDLP-related proteins as determined by molecular replacement as described above and in Example 2 below, comprise a root mean square deviation (rmsd) of no greater than 2.0 Å in the positions of Co atoms for at least 50% or more of the amino acids of the structure over the 375 residues of full-length HDLP. Such a rmsd may be expected based on the amino acid sequence identity. Chothia & Lesk, 1986, Embo J. 5:823-826.

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The refined three-dimensional HDLP structures of the present invention, specifically apo-HDLP, Cys75Ser/Cys77Ser double mutant HDLP comprising a zinc atom in the active site, HDLP/TSA complex comprising a zinc atom in the active site, and HDLP/SAHA complex comprising a zinc atom in the active site, are represented by the atomic coordinates set forth in Figures 16 to 19 respectively. The refined model for apo-HDLP comprising amino acids 1-375 consists of wild-type HDLP residues 2 to 373 with residues 1, 374 and 375 not modeled and presumed disordered and was determined to a resolution refined model for Similarly, the Å. Cys75Ser/Cys77Ser double mutant HDLP comprising a zinc atom in the active site also consists of residues 2 to 373 with residues 1, 374 and 375 not modeled and presumed disordered and was determined to a resolution of 2.0 Å. The refined model for the HDLP/TSA complex comprising a zinc atom in the active site consists of the Cys75Ser/Cys77Ser double mutant HDLP residues 2 to 373 with residues 1, 374 and 375 not modeled and presumed disordered, has TSA in the binding pocket and was determined to a resolution of 2.1 Å. HDLP/SAHA complex is similar to the HDLP/TSA complex but has SAHA in the binding pocket and was determined to a resolution of 2.5 Å.

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For the purposes of further describing the structure of HDLP and HDLP-related proteins, including, but not limited to, HDACs, from the data obtained from the HDLP crystals of the present invention, the definition of the following terms is provided:

The term " β sheet" refers to two or more polypeptide chains (or β strands) that run alongside each other and are linked in a regular manner by hydrogen bonds between the main chain C=O and N-H groups. Therefore all hydrogen bonds in a beta-sheet are between different segments of polypeptide. Most β -sheets in proteins are all-parallel (protein interiors) or all-antiparallel (one side facing solvent, the other facing the hydrophobic core). Hydrogen bonds in antiparallel sheets are perpendicular to the chain direction and spaced evenly as pairs between strands. Hydrogen bonds in parallel sheets are slanted with respect to the chain direction and spaced evenly between strands.

The term " α helix" refers to the most abundant helical conformation found in globular proteins. The average length of an α helix is 10 residues. In an α helix, all amide protons point toward the N-terminus and all carbonyl oxygens point toward the C-terminus. The repeating nature of the phi, psi pairs ensure this orientation. Hydrogen bonds within an α helix also display a repeating pattern in which the backbone C=O of residue X (wherein X refers to any amino acid) hydrogen bonds to the backbone HN of residue X+4. The α helix is a coiled structure characterized by 3.6 residues per turn, and translating along its axis 1.5 Å per amino acid. Thus the pitch is 3.6x1.5 or 5.4 Å. The screw sense of alpha helices is always right-handed.

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The term "loop" refers to any other conformation of amino acids (i.e. not a helix, strand or sheet). Additionally, a loop may contain bond interactions between amino acid side chains, but not in a repetitive, regular fashion.

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Amino acid residues in peptides shall herein after be abbreviated as follows: Phenylalanine is Phe or F; Leucine is Leu or L; Isoleucine is Ile or I; Methionine is Met or M; Valine is Val or V; Serine is Ser or S; Proline is Pro or P; Threonine is Thr or T; Alanine is Ala or A; Tyrosine is Tyr or Y; Histidine is His or H; Glutamine is Gln or Q; Asparagine is Asn or N; Lysine is Lys or K; Aspartic Acid is Asp or D; Glutamic Acid is Glu or E; Cysteine is Cys or C; Tryptophan is Trp or W; Arginine is Arg or R; and Glycine is Gly or G. For further description of amino acids, please refer to Proteins: Structure and Molecular Properties by Creighton, T.E., W.H. Freeman & Co., New York 1983.

acid having a positively charged side chain under normal physiological conditions. Examples of positively charged amino acids are Arg, Lys and His. The term "negatively charged amino acids are acid" refers to any amino acid having a negatively charged side chain under normal physiological conditions. Examples of negatively charged amino acids are Asp and Glu. The term "hydrophobic amino acid" refers to any amino acid having an uncharged, nonpolar side chain that is relatively insoluble in water. Examples of hydrophobic amino acids are Ala, Leu, Ile, Gly, Val, Pro, Phe, Trp and Met. The term "hydrophilic amino acid" refers to any amino

acid having an uncharged, polar side chain that is

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relatively soluble in water. Examples of hydrophilic amino acids are Ser, Thr, Tyr, Asp, Gln, and Cys. The term "aromatic amino acid" refers to any amino acid comprising a ring structure. Examples of aromatic amino acids are His, Phe, Trp and Tyr.

The term "charge relay system" refers to a His-Asp arrangement as described by Fersht & Sperling, 1973, J. Mol. Biol. 74:137-149; Blow et al., 1969, Nature 221:337-340.

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information obtained from The the three-dimensional structures of the present invention reveal that HDLP has a single-domain structure that belongs to the open α/β class of folds (see, e.g., Branden, 1980, Q. Rev. Biophys. 13:317-38). Two orthogonal views of the overall three-15 dimensional structure of HDLP are depicted in Figure 4A and The HDLP structure has a central eight-stranded 4B. parallel β sheet (strands arranged as $\beta 2 - \beta 1 - \beta 3 - \beta 8 - \beta 7 - \beta 4 - \beta 5 \beta6$), and sixteen α helices (labeled $\alpha1$ through $\alpha16$ 20 respectively). See Figure 4C. Four of the helices pack on either face of the β sheet (α 7, α 8, α 9, α 10 and α 11, α 12, α 13, α 14) forming the core α/β structure characteristic of this class of folds. Most of the remaining eight helices are positioned near one side of the β sheet, near stands β 2-25 β1-β3-β8. Large, well defined loops (Loops L1-L7; Figure 4C) originate from the C-terminal ends of the β -strands. The extra helices and the large L1-L7 loops are associated with a significant extension of the structure beyond the core α/β motif. This extension of the structure gives rise 30 to two prominent architectural features: a deep, narrow pocket and an internal cavity adjacent to the pocket. These

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two architectural features comprise the active site (see Figure 5A). The structure of HDLP-related proteins (e.g. HDACs) may also comprise the conserved α/β structure characteristic.

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The term "active site" comprises any or all of the following sites in HDLP, the substrate binding site, the site where the cleavage of an acetyl group from a substrate occurs or the site where an inhibitor of the HDAC family or, more particularly, HDLP binds. The active site, as referred to herein, comprises Aspl66, Asp258, His170, Tyr297, His131, His132, Aspl68, Asp173, Phe141, Phe198, Leu265, Pro22 and Gly140, and also a metal bound at the bottom of the pocket by Asp173, Aspl68 and His defined by the coordinates listed in Figures 16 to 19 with an rmsd of 2.0 Å. The metal which binds at the bottom of the pocket will be a divalent cation selected from the group consisting of zinc, cobalt or manganese.

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The deep narrow pocket has a tube-like shape with a depth of ~ 11 Å. The pocket opening constricts half way down to ~ 4.5 by 5.5 Å, and becomes wider at the bottom (see Figure 5A). The pocket and its immediate surroundings are made up of loops L1 through L7.

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The walls of the pocket are covered with side chains of hydrophobic and aromatic residues (Pro22, Tyr91 near the entrance; and Gly140, Phe141, Phe 198, Leu265 and Tyr297 further down; Figure 5B). For numbering of amino acids please refer to SEQ ID NO:1. Of particular interest are Phe141 and Phe198, whose phenyl groups face each other in

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parallel at a distance of 7.5 Å, marking the most slender portion of the pocket (see Figure 5B). Of particular interest is that only one pocket residue differs in HDAC1 when the sequences are aligned (alignment may be accomplished using DNAstar™ MegAlign™ program, Madison, WI), this residue is Glu98 of HDAC1 which is Tyr91 in HDLP. The structure reveals that this residue in HDLP is mostly solvent exposed.

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Near the bottom of the pocket of the active site at its narrowest point, is located a zinc ion (see Figure 6A). In order to obtain the zinc in the structure, the crystals may be soaked in zinc (e.g. ZnCl₂) or co-crystalized in the presence of zinc. The zinc ion is coordinated by Asp168 (Oδ1, 2.1 Å), His170 (Nδ1, 2.1 Å), Asp258 (Oδ1, 1.9 Å) and a water molecule (2.5 Å). See Figure 5B and 6B. The amino acid residues that coordinate zinc are arranged in a tetrahedral geometry, but the position of the water molecule, which is also hydrogen bonded to His131, deviates from this geometry by ~25°.

In addition to the zinc ligands, the bottom of the pocket contains two histidine (His131 and His132), two aspartic acids (Asp166 and Asp173) and a tyrosine (Tyr297). See Figure 5B and 10B. Each of the histidines makes a hydrogen bond through its No1 to an aspartic acid carboxylate oxygen, with the oxygen located in the plane of the imidizole ring (Figure 5B). This His-Asp arrangement is characteristic of the charge relay system present in the active sites of serine proteases, where it serves to polarize the imidizole Ne and increase its basicity. Fersht & Sperling, 1973, J.

Mol. Biol. 74:137-149; Blow et al., 1969, Nature 221:337-340.

The Asp166-His131 charge pair relay (hereafter referred to as "buried charged relay") is positioned even deeper in the pocket and more buried compared to the Asp173-His132 charge relay (hereafter referred to as "exposed charge relay") which is partially solvent exposed. The buried charge relay makes a hydrogen bond (2.6 Å) to the zinc-bound water molecule referred to above, and this hydrogen bond could contribute to the deviation of the water-zinc coordination from ideal geometry (Figure 5B). The exposed charge relay is directed to a point ~ 2.5 Å away from the water molecule and closer to the surface.

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Tyr 297 is positioned next to the zinc, opposite from where the two charge relay systems are located. The Tyr hydroxyl group lies 4.4 Å away from the zinc atom and has no interactions with the rest of the protein (Figure 5B). Next to Tyr297, there is an opening in the pocket wall, which leads to the adjacent internal cavity.

The floor of the internal cavity is made up of portions of the L3 and L7 loops as they emerge from the β strands, and the roof is made up by the $\alpha 1\text{-L} 1\text{-}\alpha 2$ segment. The L1 loop appears more flexible than other loops in the structure. This may allow the transient exchange of the cavity contents with the bulk solvent.

The cavity is lined primarily with hydrophobic residues and is particularly rich in glycine residues (Ala127, Gly128, Gly129, Met130, and Phe141 of L3; Gly293, Gly294, Gly295 and

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Gly296 of L7; and Tyr17, Pro22 and Leu23 of L1). There are only two charged residues in the cavity (Arg27 and His 21) and these are contributed by the L1 loop.

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The cavity may provide space for the diffusion of the acetate product away from the catalytic center, which may otherwise be crowded and shielded during deacetylation from the solvent when the substrate is bound. Such a role for the cavity is supported by the observation that the cavity contains three water and two isopropanol molecules (from the crystallization buffer) in the 1.8 Å apo-protein structure. The cavity may also bind another cofactor, in addition to zinc, for the facilitation of the enzymatic activity of the HDLP. A proposed catalytic mechanism for deacetylation is provided in Figure 8.

The structure of HDLP as defined by the present invention, in conjunction with the HDAC1 sequence homology, shows that the 375-amino acid HDLP protein corresponds to the histone deacetylase catalytic core which is conserved across the HDAC family (see Figure 2). The 35.2% HDLP-HDAC1 sequence identity predicts structural similarity with a rmsd in $C\alpha$ positions of ~ 1.5 Å. Chothia and Lesk describe the relation between the divergence of sequence and the structure of proteins in Embo J. 5:823-826 (1986). The 40residue C-terminus of HDLP is likely to have a divergent structure since this region has lower homology to HDAC1, although the α 16 helix in this region is part of the conserved open α/β core fold and HDAC1 is likely to comprise a similar helix. However divergent this C-terminal region may be, this region is outside the active site and is likely to not effect the structure of the active site. Beyond the

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C-terminus of the histone deacetylase catalytic core, HDAC family members are divergent in length and sequence. In the HDAC family, this region (amino acid residues ~390-482) is highly polar, populated with acidic residues, and is likely to be flexible or loosely folded.

The HDLP-HDAC homology maps primarily to the hydrophobic core and to the L1-L7 loops, with portions of the loops that make up the pocket and adjacent cavity having the highest level of amino acid residue sequence conservation (Figure 9A and 9B). Specifically, all of the polar residues in the active site (the zinc ligands, the two charge relay systems, and Tyr297) and the hydrophobic residues that make up the walls of the pocket (Gly140, Phe141, Phe198 and Leu265) are identical. Among the residues that make up the internal cavity, the ones closest to the active site are either identical or conservatively substituted (for example, Leu23 \rightarrow Met and Met130 \rightarrow Leu). Surface residues around the pocket are conserved to a lesser extent, but are still above 35% average sequence identity.

The information obtained from the inhibitor-bound HDLP complex crystal structures of the present invention reveal detailed information which is useful in the design, isolation, screening and determination of potential inhibitor compounds which may inhibit HDLP/HDAC family members. As described above, the HDLP structure consists of a parallel β sheet with α helices packing against both faces (Figure 4A, 4B, and 4C). At one end of the β sheet, 7 loops (L1-L7) form a narrow, tube-like pocket which are lined with hydrophobic residues and which comprise a zinc binding site, several polar side chains, including two Asp-His charge

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relay systems. Mutation of the zinc ligands and other polar residues at the pocket bottom reduces or eliminates the catalytic activity.

The present inventors found that mutation at the Tyr297Phe site reduced activity. See also, Hassig et al., 1998, Proc. Natl. Acad. Sci. USA 95:3519-3524; Kadosh & Struhl, 1998, Genes Dev. 12:797-805. The elimination of activity by mutation of these residues indicates that this region is the enzyme active site. Adjacent to the active site, there is an internal cavity that may provide space for the diffusion of the acetate reaction product. Homology at the active site between HDLP and HDAC1, as described above, indicates that they share structural and functional homology.

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The inhibitor compound, trichostatin A (TSA) (Tsuji et al., 1976, J. Antibiotics 29:1-6) binds HDLP by inserting its long aliphatic chain, which has a hydroxamic acid group at one end, into the pocket (Figure 6A, 6B and 6C). aliphatic chain makes multiple contacts in the well-like, hydrophobic portion of the pocket. The hydroxamic acid reaches the polar bottom of the pocket, where it coordinates the zinc in a bidentate fashion and also forms hydrogen bonds with the polar residues in the active site, including the two charge relay system histidines. dimethylamino-phenyl group at the other end of the TSA chain makes contacts at the pocket entrance and serves to cap it. The amino acid residues of HDLP which contact TSA are conserved in HDAC, indicating that TSA binds and inhibits HDAC in a similar fashion to HDLP.

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In the complex, the hydroxamic acid, most of the aliphatic chain and part of the dimethylamino-phenyl group of TSA are buried (60% of TSA's surface area; Figure 6A). The hydroxamic acid group binds the zinc in a bidentite fashion forming bonds through its carbonyl (2.4 Å) and hydroxyl groups (2.2 Å) resulting in a penta-coordinated Zn²+ (Figure 6B and 6C). The hydroxamic acid hydroxyl group replaces the water molecule that binds to the zinc in the apo-HDLP structure described above. The hydroxamic acid also hydrogen bonds with both charge relay system histidines (hydroxyl oxygen to His131 Ne2, 2.8 Å; and nitrogen to His132 Ne2, 2.8 Å), and the Tyr297 hydroxyl group (2.4 Å; Figure 6B and 6C).

The 5-carbon long branched alkene chain of TSA fits snugly in the narrow portion of the pocket making multiple van der Waals contacts with all of the hydrophobic groups lining the pocket (Figure 6B and 6C). Near its center, the chain contains a methyl substituted carbon-carbon double bond which is sandwiched between the phenyl groups of the Phel41 and Phe98 at the tightest point of the pocket (Figure 6A and 6B). The length of the alkene chain appears optimal for spanning the length of the pocket, and allowing contacts both at the bottom and at the entrance of the pocket, although, the cap group of Formula (I) may provide length to span the pocket allowing for a shorter alkene chain (aliphatic chain).

At the entrance of the pocket, one face of the planar structure formed by the dimethylamino-phenyl and adjacent carbonyl groups of TSA makes contacts at the rim of the pocket (Pro22, Tyr91, Phe141; Figure 6B and 6C). This

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packing is facilitated by the roughly 110° angle in the overall structure of TSA at the junction of the aliphatic chain and the dimethylamino-phenyl group (occurring at the sp³ hybridized C8 carbon). Upon TSA binding, the side chain of Tyr91, which is mostly solvent exposed, changes conformation to make space for the dimethylamino-phenyl group. This is the only change near the active site observed upon TSA binding.

10 The hydroxamic acid group is a common motif in zinc metalloprotease inhibitors. See U.S. Patent No. 5,919,940 and 5,917,090; See also, Grams et al., 1995, Biochemistry 34:14012-14020; Lovejoy et al., 1999, Nat. Struct. Biol. 6:217-221; and Holmes & Matthews, 1981, Biochemistry 15 20:6912-6920. Like TSA, these inhibitors also coordinate the active site zinc in a bidentate fashion using their hydroxamate hyroxyl and carbonyl oxygens, replace the nucleophilic water molecule with their hydroxamate hydroxyl groups and form hydrogen bonds to the general base (Grams et al., 1995, Biochemistry 34:14012-14020; Lovejoy et al., 20 1999, Nat. Struct. Biol. 6:217-221; and Holmes & Matthews, 1981, Biochemistry 20:6912-6920).

SAHA, which has a ~30-fold weaker inhibitory activity than

TSA (Richon et al., 1998, Proc. Natl. Acad. Sci. USA

95:3003-3007), binds HDLP similarly to TSA (see, e.g.,

Figure 4D). The SAHA hydroxamic acid group makes the same

contacts to the zinc and active site residues, and the

importance of these interactions is underscored by the loss

of activity of SAHA derivatives lacking the hydroxamic group

(Richon et al., 1998, Proc. Natl. Acad. Sci. USA 95:3003-

3007). The six-carbon long aliphatic chain of SAHA packs in the tube-like hydrophobic portion of the pocket. Compared to TSA however, SAHA's aliphatic chain packs less snugly and makes fewer van der waals contacts, in part, because SAHA lacks TSA's C15 methyl group branch. SAHA also lacks TSA's double bonds in this region, and this may lead to increased flexibility of the aliphatic chain. The cap group of SAHA consists of a phenyl-amino ketone group. In the crystal structure, the phenyl group has weak electron density, suggesting that it does not pack as well as the cap group of TSA. This may be due to the larger separation between the hydroxamic and cap groups of SAHA compared to TSA (compare TSA, Formula (II) and SAHA, Formula (III), below).

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25 (III)

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The determination of the structure of HDLP and HDLP bound to an inhibitory compound has enabled, for the first time, the identification of the active site of HDLP and of related HDLP proteins, such as proteins belonging to the HDAC family.

The three-dimensional structural information and the atomic coordinates associated with said structural information of HDLP bound to an inhibitory compound is useful in rational drug design providing for a method of identifying inhibitory compounds which bind to and inhibit the enzymatic activity of HDLP, HDAC family proteins and other histone deacetylaselike proteins related to HDLP. Said method for identifying said potential inhibitor for an enzyme comprising deacetylase activity comprises the steps of (a) using a three-dimensional structure of HDLP as defined by its atomic coordinates listed in Figure 16 to 19; (b) employing said three-dimensional structure to design or select said potential inhibitor; (c) synthesizing said potential inhibitor; (d) contacting said potential inhibitor with said enzyme in the presence of an acetylated substrate; and (e) determining the ability of said inhibitor to inhibit said deacetylase activity.

The potential HDLP and HDLP-related (e.g. HDAC) inhibitors identified by the method of the present invention are represented by formula (I)

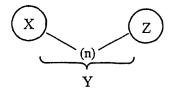
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(I)



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wherein X comprises a cap group which binds to at least one amino acid selected from the group consisting of proline and leucine; Y comprises an aliphatic chain group which binds to at least one amino acid selected from the group consisting of leucine, phenylalanine and glycine; and Z comprises an active site binding group which binds to at least one amino acid selected from the group consisting of aspartic acid, tyrosine and histidine and wherein Z may further bind to a zinc atom and with the provision that the compound of Formula (I) is not TSA, trapoxin, SAHA, SAHA derivatives described in U.S. Patent Nos. 5,608,108; 5,700,811; 5,773,474; 5840,960 and 5,668,179.

The present invention permits the use of molecular design 20 techniques to design, identify and synthesize chemical entities and compounds, including inhibitory compounds, capable of binding to the active site of HDLP and HDLPrelated proteins. The atomic coordinates of apo-HDLP and inhibitor-bound HDLP may be used in conjunction with 25 computer modeling using a docking program such as GRAM, DOCK, HOOK or AUTODOCK (Dunbrack et al., 1997, Folding & Design 2:27-42) to identify potential inhibitors of HDLP and HDLP-related proteins (e.g. HDAC1). This procedure can 30 include computer fitting of potential inhibitors to the

active site of HDLP to ascertain how well the shape and the

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the potential inhibitor will chemical structure of complement the active site or to compare the potential inhibitors with the binding of TSA or SAHA in the active See Bugg et al, 1998, Scientific American December: 92-98; West et al., 1995, TIPS 16:67-74. potential inhibitors designed by modeling with a docking program conform to the general formula (I) as described Computer programs may also be employed to estimate the attraction, repulsion and stearic hindrance of the HDLP and potential inhibitor compound. Generally, the tighter the fit, the lower the stearic hindrances, the greater the attractive forces, and the greater the specificity which are important features for a specific inhibitory compound which is more likely to interact with HDLP and HDLP-related proteins rather than other classes of proteins. features are desired particularly where the inhibitory compound is a potential antitumor drug.

The compounds of the present invention may also be designed by visually inspecting the three-dimensional structure to determine more effective deacetylase inhibitors. This type of modeling may be referred to as "manual" drug design. Manual drug design may employ visual inspection and analysis using a graphics visualization program such as "O" (Jones, T.A., Zhou, J.Y., Cowan, S.W., and Kjeldgaard, M., Improved method for building protein models in electron density maps and the location of errors in these models, Acta Crystallog., A47, 110-119.

Initially potential inhibitor compounds can be selected for their structural similarity to the X, Y and Z constituents

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of formula (I) by manual drug design. The structural analog thus designed can then be modified by computer modeling programs to better define the most likely effective candidates. Reduction of the number of potential candidates is useful as it may not be possible to synthesize and screen a countless number of variations compounds that may have some similarity to known inhibitory molecules. Such analysis has been shown effective in the development of HIV protease inhibitors (Lam et al., 1994, Science 263:380-384; Wlodawer et al., 1993, Ann. Rev. Biochem. 62:543-585; Appelt, 1993 Perspectives in Drug Discovery and Design 1:23-48; Erickson, 1993, Perspectives in Drug Discovery and Design 1:109-128. Alternatively, random screening of an small molecule library could lead to potential inhibitors whose inhibitory activity may then be analyzed by computer modeling as described above to better determine their effectiveness as inhibitors.

The compounds designed using the information of the present invention may be competitive or noncompetitive inhibitors. These designed inhibitors may bind to all or a portion of the active site of HDLP and may be more potent, more specific, less toxic and more effective than known inhibitors for HDLP and HDLP-related proteins, particularly HDACs. The designed inhibitors may also be less potent but have a longer half life in vivo and/or in vitro and therefore be more effective at inhibiting histone deacetylase activity in vivo and/or in vivo for prolonged periods of time. Said designed inhibitors are useful to inhibit the histone deacetylase activity of HDLP and HDLPrelated proteins (e.g. HDAC1), to inhibit cell growth in

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vitro and in vivo and may be particularly useful as antitumor agents.

The present invention also permits the use of molecular design techniques to computationally screen small molecule data bases for chemical entities or compounds that can bind to HDLP in a manner analogous to the TSA and SAHA as defined by the structure of the present invention. Such computational screening may identify various groups which may be defined as "X", "Y" or "Z" of formula (I) above and may be employed to synthesize the potential inhibitors of the present invention comprising formula (I). potential inhibitors may be assayed for histone deacetylase inhibitory activity in a histone deacetylase activity assay (see Example 3 below), may be co-crystallized with HDLP to determine the binding characteristics through X-ray crystallography techniques defined above (e.g. said cocrystal structure may be determined by molecular replacement to assess the binding characteristics of said potential inhibitor), or may be assessed based on binding activity by incubating said potential inhibitor with said HDLP, performing gel filtration to separate any free potential inhibitor to HDLP-bound inhibitor, and determining the amount of histone deacetylase activity of the inhibitorbound HDLP. To measure binding constants (e.g., Kd), methods known to those in the art may be employed such as Biacore™ analysis, isothermal titration calorimetry, Elisa with a known drug on the plate to show competitive binding, or by a deacetylase activity assay.

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The design of potential inhibitors of the present invention is further facilitated by reference to Figure 9, which is a surface representation figure that depicts the surface grooves. Analysis of such grooves gives insight into the constituents of the cap group of formula (I). The surface grooves are labeled groove A, groove A', groove B and groove C, into which additional cap groups may bind. The structure of HDLP bound to either TSA or SAHA shows that the cap groups of TSA and SAHA bind in groove A. By analysis of the amino acid sequence identity of HDLP and HDACs, Groove A is well conserved in HDACs, has a significant hydrophobic component, appears deep enough to allow for significant interactions and is also the largest of the four grooves. In addition to the dimethylamino phenyl group of the TSA, the A groove can fit approximately 200 daltons worth of groups (e.g. groove A could accommodate a naphthalene-like group after an appropriate spacer, etc.). Groove A, as referred to herein, is characterized by the following conserved residues of HDLP: His 21, Pro22, Lys24, Phe141, The periphery of groove A comprises Leu265 and Phe335. unconserved residues. Additionally, Groove A', as referred to herein, comprises primarily unconserved residues.

Groove B is immediately adjacent to the pocket. Of significance is that the bottom of groove B comprises the N-epsilon nitrogen of His170, which coordinates the zinc through its N-delta nitrogen. Significant binding energy may be achieved by contacting the Ne proton of His170 with a carboxylic acid or sulfate group. In addition, groove B may be large enough to fit a phenyl group, the face of which may comprise a partial negative charge which may pack over the N-epsilon proton of His170. The conserved residues of

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groove B, as referred to herein are: His170, Tyr196 and Leu265.

Groove C is not as well conserved as the other two grooves and the amino acid residues which comprise groove C are mostly polar and solvent exposed. Groove C, as referred to herein comprises the following conserved residues: Asn87, Gly140 and Phe198.

The compounds of the present invention are represented by formula (I):

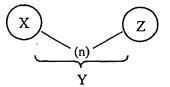
(I)

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Examples for suitable X constituents wherein X comprises a cap group may be described in three categories, depending upon which surface of groove A, A', B and/or C they are targeted to. The cap group may comprise all three categories on the same compound. Of particular benefit may be replacing the cap group of TSA or SAHA with a large, rigid structure. Nonlimiting examples for suitable cap groups (X) of formula (I) which may bind in groove A are: (1) attaching a 1-3 methyl linker followed by a phenyl or naphthalene group from the para or meta position of SAHA's phenyl group represented by formula (IV):

- 45 -

(IV)

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(2) attaching a 2-3 methyl linker followed by a phenyl or naphthalene group from the meta position of TSA's phenyl cap group, or from TSA's dimethyl amino group represented by formula (V):

(V)

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and which may bind in groove B is a 1-3 methyl group spacer followed by a carboxylate, sulfate or phenyl group as represented by formula (VI):

(VI)

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With respect to the aliphatic (Y) group, the diameter of the pocket suggests that one more methyl "side chain" could fit, in addition to the C15 methyl group on the C10 carbon. Nonlimiting suitable examples for Y constituents wherein Y comprises an aliphatic chain group are as follows: (1) add

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a methyl group to TSA on the C12 carbon (with or without a methyl group on the C10 carbon and with or without double bonds and with or without substituting the X and/or Z constituents of formula (I)as represented by formula (VII):

5 (VII)

10 (2) add a methyl group to TSA on the C9 carbon (with or without a methyl group on the C10 carbon; with or without both or either of the double bonds, and with or without substituting the X and/or Z constituents of formula (I) as represented by formula (VIII):

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(VIII)

$$\begin{array}{c|c} X & \begin{pmatrix} 14 & \begin{pmatrix} 15 \end{pmatrix} & \begin{pmatrix} 15 \end{pmatrix} \\ & 1 \end{pmatrix} & \begin{pmatrix} 15 \end{pmatrix} & \end{pmatrix} & \begin{pmatrix} 15 \end{pmatrix}$$

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(3) replace the two alkalene double bonds of TSA with only one between C10 and C11, which may free the C11 and C12 torsion to allow for a better fit, the X and/or Z groups may also be substituted as represented by formula (IX):

25 (IX)

(4) cyclize C15 and C12 carbons of TSA through a sulphur atom (or nitrogen atom), the X and/or Z groups may also be substituted as represented by formula (X):

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(X)

(5) extend from the C9 carbon of TSA such that the extension approaches and/or enters groove B (see Figure 9); making C9 sp3 so that it can have some freedom; attach to C9 a 1-3 methyl group spacer which may include a double bond and they attaching thereto a sulfate, carboxylate, sulfate, hyroxyl, or phenyl group which may make an interaction with the N-epsilon proton of His170 which may coordinate the zinc atom as represented by formula (XI):

(XI)

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$$\begin{array}{c|c} X & \begin{array}{c} & & & \\ & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\ & & \\$$

25 (6) extend off the C8 carbon (replacing C14) of TSA such that the extension approaches or enters groove B; attach a 1-3 methyl group spacer (which may include a double bond) and then link thereto a carboxylate, sulfate, hydroxyl or phenyl group such that an interaction is made with the N-epsilon proton of His170 that coordinates the zinc atom; the X and/or Z constituents may also be substituted as represented by formula (XII):

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(XII)

$$\begin{array}{c|c} R_1 \\ (CH_2)n & (_{15}) \\ \hline \\ X \end{array} \begin{array}{c} (CH_2)n & (_{15}) \\ \hline \\ R_1 = \begin{cases} -COOH \\ -SO_4 \\ -OH \end{cases} \end{array}$$

(7) substitute the C8 carbon at the end of the aliphatic chain such that the substitution may contact groove A, A', B and or C, in such an example, a cap group (X) may or may not be required and the X and Z constituents may be substituted as well, as represented by formula (XIII):

(XIII)

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(8) formulas VII through XIII above wherein the aliphatic chain further comprises a methyl group between the active site binding group (Z) and the C8 carbon, and preferably just before the C8 carbon, increasing the distance between X and Z, (9) make the connection between the aliphatic chain and the cap group more rigid (e.g., by closing a 6-membered ring which may or may not comprise oxygen, the X and Z group may also be substituted as represented by formula (XIV):

(XIV)

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and (10) combining two or more of the changes depicted by formulas (VII-XIV).

Additionally, nonlimiting examples for suitable Z groups wherein Z comprises an active site binding group are as follows: (1) hydroxamic acid, (2) carboxylic acid, (3) sulfonamide, (4) acetamide, (5) epoxyketone, (6) an ester with a methyl linker and a hydroxyl of acetate ester group to lead into the cavity and interact with a conserved arginine (Arg27) as represented by formula (XV): (VV)

$$R_1$$
 CH_2 CH_2 CH_2 CH_3 CH_4 CH_4 CH_5 CH_5 CH_6 C

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and (7) an alphaketone as represented by formula (XVI): (XVI)

$$\frac{HQ}{C - G} = \begin{cases} - - (CH_2)n - OH \\ - - (CH_2)n - C \end{cases}$$

$$\frac{Q}{Q} = \begin{cases} - - (CH_2)n - OH \\ - - (CH_2)n - C \end{cases}$$

$$\frac{Q}{Q} = \begin{cases} - - (CH_2)n - OH \\ - - (CH_2)n - C \end{cases}$$

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Additionally, other suitable X, Y and Z constituents may be the skilled artisan given the threeenvisioned by dimensional structural information of the present invention.

After having determined potential suitable X, Y and Z constituents, the constituents are combined to form a compound of formula (I) using combinatorial chemistry techniques. This may be achieved according to U.S. Patent 5,608,108; 5,700,811; 5,773,474; 5,840,960 5,668,179, incorporated herein by reference. Any methods

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known to one of skill in the art may be employed to synthesize compounds of formula (I) comprising X, Y and Z constituents as determined by the methods described above.

As mentioned above, the compounds of formula (I) are useful to inhibit the histone deacetylase activity of HDLP and HDAC-related proteins. Such inhibition may allow for a reduction or cessation of cell growth in vitro and in vivo.

For in vitro use, such reduction or cessation of cell growth is useful to study the role of histone deacetylation and differentiation during the cell cycle and also to study other mechanisms associated with cell cycle arrest and particularly how the repression of transcription is involved in cell cycle progression which may be studies in a yeast model system such as that described by Kadosh & Struhl, 1998, Mol. Cell. Biol. 18:5121-5127. In vitro model systems which may be employed to study the effects of potential inhibitors on cell cycle progression and also tumor growth include those described by: Richon et al., 1998, Proc. Natl. Acad. Sci. USA 95:3003-3007; Yoshida et al., 1995, Bioessays 17:423-430; Kim et al., 1999, Oncogene 18:2461-2470; Richon et al., 1996, Proc. Natl. Acad. Sci. USA 93:5705-5708; and Yoshida et al., 1987, Cancer Res. 47:3688-3691.

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For in vivo use, such a reduction or cessation of cell growth is useful to study the effect of said inhibitor compounds in non-human animal model systems of cancer and is also useful for the treatment of cancer in a recipient in need of such treatment. Non-limiting examples of animals which may serve as non-human animal model systems include

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mice, rats, rabbits, chickens, sheep, goats, cows, pigs, and non-human primates. See, e.g., Desai et al., 1999, Proc. AACR 40: abstract #2396; Cohen et al., 1999, Cancer Res., submitted. The compounds of the present invention may be administered to a transgenic non-human animal wherein said animal has developed cancer such as those animal models in which the animal has a propensity for developing cancer (e.g. animal model systems described in U.S. Patents 5,777,193, 5,811,634, 5,709,844, 5,698,764, and 5,550,316). Such animal model systems may allow for the determination of toxicity and tumor reduction effectiveness of the compounds of the present invention.

A preferred compound of the present invention may comprise high specific activity for HDLP and HDAC-related proteins, good bioavailability when administered orally, activity in reducing or ceasing cell growth in tumor cell lines, and activity in reducing or ceasing tumor growth in animal models of various cancers.

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Accordingly, another aspect of this invention is a method of eradicating or managing cancer in a recipient, which may be an animal and is preferably a human. Said method comprises administering to said recipient a tumor reducing amount of a compound as defined by formula (I) above, or a physiological acceptable salt thereof.

In a further aspect of the invention, there is provided a composition comprising the compound of formula (I) and an excipient or carrier. Administration of the foregoing agents may be local or systemic. Such carriers include any

suitable physiological solutions or dispersant or the like. The physiologic solutions include any acceptable solution or dispersion media, such as saline, or buffered saline. The carrier may also include antibacterial and antifungal agents, isotonic and absorption delaying agents, and the like. Except insofar as any conventional media, carrier or agent is incompatible with the active ingredient, its use in the compositions is contemplated.

10 Routes of administration for the compositions containing the delivery vehicle constructs of the present invention include any conventional and physiologically acceptable routes, such as, for example, oral, pulmonary, parenteral (intramuscular, (IV) intraperitoneal, intravenous orsubcutaneous injection), inhalation (via a fine powder formulation or a 15 fine mist), transdermal, nasal, vaginal, rectal, or sublingual routes of administration and can be formulated in dosage forms appropriate for each route of administration.

The following examples are provided to more clearly illustrate the aspects of the invention and are not intended to limit the scope of the invention.

EXAMPLES

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25 Example 1: Protein Production and Purification:

Full-length wild-type HDLP (Genbank accession number AE000719) was subcloned from an Aquifex aeolicus chromosomal DNA preparation (provided by Robert Huber of Universitaet of Regensburg, Germany) into the pGEX4T3 (Amersham-Pharmacia, Piscataway, NJ) vector using the polymerase chain reaction (PCR). The cysteine-to-serine and active site mutants were constructed by PCR site directed mutagenesis and were

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sequenced. The HDLP-qlutathione S-transferase (GST) fusion protein was produced in Escherichia coli, purified by chromatography affinity using a column qlutathione-sepharose resin (Amersham-Pharmacia, Piscataway. NJ), and by anion-exchange chromatography (Q-sepharose™; Amersham-Pharmacia, Piscataway, NJ). HDLP was cleaved from the fusion protein with thrombin at 4° C, was purified by (Q-sepharose™; anion-exchange Amersham-Pharmacia, Piscataway, NJ) and gel filtration chromatography (Superdex™200; Amersham-Pharmacia, Piscataway, NJ), and was concentrated to typically 25 mg/ml in a buffer of 25 mM bis-tris propane (BTP), 500 mM NaCl, 5 mM dithiothrietiol (DTT), 2% isopropanol, pH 7.0.

15 Although, it is not known what metal cofactor HDLP contains in vivo, it is presumed to be zinc because of the arrangement of the ligands and the similarities in the active site to the zinc proteases. The lack of metal in the purified HDLP is presumed due, in part, to the use of DTT 20 during purification. HDLP was reconstituted with Zn2+ by mixing the Cys75Ser/Cys77Ser double mutant at 10 mg/ml with a 5-fold molar excess of ZnCl2 in a buffer of 25 mM bis-tris propane, 200 mM NaCl, 1% isopropanol, pH 7.0. Unbound ZnCl² was removed by fractionating HDLP through a G25 desalting 25 column (Amersham-Pharmacia, Piscataway, NJ). The HDLP-Zn2+-TSA complex was prepared by incubating the Zn2+ reconstituted HDLP mutant with 1 mM TSA for 45 minutes, followed by gel filtration chromatography (Superdex™200; Amersham-Pharmacia, Piscataway, NJ) to remove excess TSA, 30 and concentration to typically 25 mg/ml in a buffer of 25 mM bis-tris propane, 500 mM NaCl, 1% isopropanol, pH 7.0.

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FLAG epitope tagged human HDAC1 was overexpressed using a baculovirus expression system in Hi5 (Invitrogen, Carlsbad, CA) insect cells grown in suspension in serum-free media (Sf900, Gibco, Grand Island, NY). The fusion protein was purified by anion exchange and affinity chromatography using Anti-FLAG M2 affinity resin (Sigma, St. Louis, MO) and FLAG Peptide (Sigma,, St. Louis, MO).

Example 2: Crystallization and data collection:

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10 Crystals of apo-HDLP were grown at room temperature by the hanging-drop vapor-diffusion method, from 7.5% isopropanol, 28% PEG 1500, 425 mM NaCl, 100 mM Tris-Cl, pH 7.0. They form in space group C2 with a = 51.4 Å, b = 93.8 Å, c = 78.7 Å, β = 96.9 Å, and contain one HDLP molecule in the asymmetric unit. Diffraction data were collected with crystals flash-frozen in a buffer of 7.5% isopropanol, 35% PEG 1500, 75 mM NaCl, 100 mM Tris-Cl, pH 8.0, at -170° C.

The structure of the HDLP- Zn²⁺ complex was determined from HDLP Cys75Ser/Cys77Ser double mutant crystals grown from 23% tert-butanol, 27% PEG 1500, 400 mM KCl, 100 mM bis-tris propane-Cl, pH 6.8. Space group and cell dimensions were identical to the apocrystals. The HDLP-Zn²⁺ crystals were harvested and frozen in 27% tert-butanol, 22% PEG 1500, 50 mM KCl, 20 mM NaCl, 0.2 mM ZnCl₂, 100 mM bis-tris propane, pH 6.8, at -170° C.

Crystals of the HDLP-Zn²⁺-TSA complex comprised HDLP Cys75Ser/Cys77Ser double mutant and were grown from 23% tert-butanol, 27% PEG 1500, 600 mM KCl, 100 mM bis-tris propane-Cl, pH 6.8, by microseeding. The crystals were grown in the presence of zinc. They form in space group

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 $P2_12_12_1$ with a = 53.4 Å, b = 94.4 Å, c = 156.3 Å and contain two HDLP- Zn^{2+} -TSA complexes in the asymmetric unit. The HDLP- Zn^{2+} -TSA crystals were harvested and frozen in the same cryobuffer as the HDLP- Zn^{2+} crystals except that 0.5mM TSA was added. Data were processed with DENZO and SCALEPACK (Otwinowski & Minor, 1997, Method. Ensemble. 276:307-326). MIR analysis, model building and refinement.

The HDLP-Zn²⁺-SAHA complex crystals were grown and evaluated the same as the HDLP-Zn²⁺-TSA crystals. However, the restraints for the SAHA structure were constructed based on stereochemical parameters from TSA. Like the apo-HDLP crystals, the SAHA/HDLP co-crystals grew in space group C2.

15 Heavy-atom soaks were performed with the apo-HDLP crystals in a buffer of 7.5% isopropanol, 30% PEG 1500, 75 mM NaCl, 100 mM Tris-Cl, pH 8.0, supplemented with 1.0 mM thimerosal for 2h, 5 mM KAu(CN)₂ for 1h, and 1 mM Pb(Me)₃OAc for 2h. MIR phases were calculated with the program MLPHARE (The 20 CCP4 suite: Programs for computational crystallography, 1994, Acta Crystallogr. D 50:760-763) at 2.5 Å using the anomalous diffraction signal from the thimerosal derivative, and had a mean figure of merit of 0.55. The phases were improved by solvent flattening with the program DM (The CCP4 suite: Programs for computational crystallography, 1994, 25 Acta Crystallogr. D 50:760-763) , and were used to build the initial model with the program O (Jones et al., 1991, Acta Crystallogr. A 47:110-109). Successive rounds of rebuilding and simulated annealing refinement with the program CNS (Brunger et al., 1998, Acta Crystallogr. D 30 54:905-921) allowed interpretation of HDLP from residues 2

to 373. Residues 1, 374, and 375 were not modeled and are presumed to be disordered.

The structure of the HDLP-Zn2+-TSA and HDLP-Zn2+-SAHA complex were determined by molecular replacement with the program AMORE (The CCP4 suite: Programs for computational crystallography, 1994, Acta Crystallogr. D 50:760-763) using the apo-HDLP structure as a search model. The initial electron density maps had strong and continuous difference density for the entire TSA molecule. However the SAHA molecule was not as well ordered in the cap group region. The structure of TSA was obtained from the Cambridge Structural Database (Refcode TRCHST) and was used to define stereochemical restraints used in the refinement with the program CNS. The restraints of SAHA were constructed based on stereochemical parameters from TSA and surrounding amino acid residues. The dimer interface in the HDLP-Zn2+-TSA and HDLP-Zn2+-SAHA crystals primarily involves Phe200 on the protein surface. The Phe200 side chain contacts Tyr91, whose side chain conformation changes on TSA binding, and part of the dimethyl amino phenyl group of TSA from the second protomer. The HDAC family does not contain a phenylalanine residue at the equivalent position.

25 Example 3: Histone deacetylase assays:

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Purified proteins were assayed by incubating 10 μ g of [³H]acetyl-labeled murine erythroleukemia histone substrate and HDAC assay buffer (20 mM Tris-HCl, pH 8.0, 150 mM NaCl, 10% glycerol) for 30-60 minutes at 37° C in a total volume of 30 μ l. The final concentrations of HDLP and HDACl-FLAG were 3.6 μ M and 0.24 μ M, respectively. Assays were performed in duplicate. The reactions were stopped and the

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released acetate was extracted and assayed as described (Hendzel et al., 1991, J. Biol. Chem. 266:21936-21942). [3H] acetyl-labeled murine erythroleukemia histones were prepared essentially as described (Carmen et al., 1996, J. Biol. Chem. 271:15837-15844). Inhibitors were added in the absence of substrate and incubated on ice for 20 minutes, substrate was added, and the assay performed as described above. HDLP was inclubated with 20 μ M ZnCl₂ and 20 μ M MnCl₂(H2O)₄ in HDAC buffer and tested for activity.

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Only HDLP dialyzed against ${\rm ZnCl_2}$ had activity. HDAC1-FLAG was dialyzed against 20 μM ${\rm ZnCl_2}$ in HDAC buffer which had no effect on activity. Therefore, HDAC1-FLAG contains a metal as purified.

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The *in vivo* substrate of HDLP is not known. HDLP may have a role in acetoin utilization like the *B. subtilis* AcuC gene product, and it has been annotated as such in the genome sequence, but the reaction catalyzed by AcuC is also not known. Furthermore, the *A. aeolicus* genome appears to lack the acuA and acuB genes that are part of the acuABC operon of B. subtilis (Deckert et al., 1998 Nature 392:353-358), and HDLP is as similar to human HDAC1 (35.2 % identity) as it is to B. subtilis AcuC (34.7 % identity).

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What is claimed is:

- 1. A crystal of an enzyme comprising deacetylase activity wherein said crystal effectively diffracts X-rays for the determination of the atomic coordinates of said enzyme to a resolution of greater than 4 Å and wherein the structure of said enzyme comprises a conserved core α/β structure characteristic fold wherein said conserved α/β fold comprises an eight-stranded parallel β sheet and eight α helices and wherein four of the helices pack on either face of said parallel β sheet and wherein said structure of said enzyme comprises an rmsd of less than or equal to 1.5 Å in the positions of Cα atoms for at least 2/3 or more of the amino acids of HDLP as defined by the atomic coordinates of HDLP.
 - 2. The crystal of claim 1, wherein said protein structure further comprises:
 - (a) eight α helices positioned near one side of the β sheet; and
 - (b) at least seven large, well defined loops originating from the C-terminal ends of the β -strands of said eight-stranded parallel β sheet wherein the eight extra helices and the seven large loops are associated with a significant extension of the structure beyond the core α/β motif and wherein said extension of the structure gives rise to a deep, narrow pocket and an internal cavity adjacent to the pocket.

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3. The crystal of claim 1, wherein said enzyme comprising deacetylase activity is selected from the group

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consisting of HDLP, HDLP-related proteins, HDAC1, HDAC2, HDAC3, HDAC4, HDAC5, HDAC6, HDAC-related proteins, APAH, AcuC, and functional derivatives thereof.

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- 4. The crystal of claim 2 further comprising a specifically bound zinc atom in the active site of said enzyme.
- 10 5. The crystal of claim 2 further comprising a specifically bound deacetylase inhibitor compound in the active site of said enzyme.
- 6. The crystal of claim 2 define by the atomic coordinates according to Figure 16.
 - 7. A method for identifying a potential deacetylase inhibitor compound for an enzyme which comprises deacetylase activity, said method comprising the steps of:
 - a. using a three-dimensional structure of HDLP as defined by atomic coordinates according to Figure 16;
 - b. employing said three-dimensional structure to design or select said potential inhibitor;
 - c. synthesizing said potential inhibitor;
 - d. contacting said potential inhibitor with said enzyme in the presence of an acetylated substrate; and
- 30 e. determining the deacetylase inhibitory activity of said potential inhibitor.

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- 8. The method of claim 7, wherein the three-dimensional structure is designed or selected using computer modeling.
- 5 9. The method of claim 7, wherein the potential deacetylase inhibitor is designed de novo.
- 10. The method of claim 7, wherein the potential deacetylase inhibitor is designed based on a known inhibitor.
 - 11. The method of claim 7, wherein said enzyme comprising deacetylase activity is selected from the group consisting of HDLP, HDLP-related proteins, HDAC1, HDAC2, HDAC3, HDAC4, HDAC5, HDAC6, HDAC-related proteins, APAH, and AcuC.

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- 12. A method of evaluating the binding properties of the potential deacetylase inhibitor compound comprising the steps of:
 - a. co-crystallizing said compound with HDLP;
 - b. determining the three-dimensional structure of said HDLP-potential inhibitor complex co-crystal by molecular replacement using the threedimensional structure of HDLP as defined by atomic coordinates according to Figure 16; and
 - c. analyzing said three-dimensional structure of said HDLP bound to said potential inhibitor compound to evaluate the binding characteristics of said potential inhibitor compound.
 - 13. A method for solving the structure of an HDAC family

member crystal comprising the steps of:

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- a. collecting X-ray diffraction data of said crystal wherein said data diffracts to a high resolution limit of greater than 4 Å;
- b. using the atomic coordinates of HDLP accoding to Figure 16 to perform molecular replacement or refinement and difference fourier with said X-ray diffraction data of said HDAC family member crystal to determine the structure of said HDAC family member; and
 - c. refining said structure of said HDAC family member.
- 14. The method of claim 13, wherein said HDAC family member is HDAC1.
 - 15. A Cys75Ser/Cys77Ser double mutant of HDLP wherein said mutant is encoded by the nucleic acid sequence of SEQ ID NO:4.
 - 16. A Cys75Ser/Cys77Ser double mutant of HDLP wherein said mutant has the amino acid sequence of SEQ ID NO:3.
 - 17. A nucleotide sequence according to SEQ ID NO:4
 - 18. An expression vector comprising the nucleotide sequence of claim 17.
- 19. A method of using the crystal of claim 1 for screening for a novel drug comprising:
 - a. selecting a potential ligand by performing

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rational drug design with the three-dimensional structure determined for the crystal;

contacting the potential ligand with the ligand
 binding domain of the crystal; and

c. detecting the binding potential of the potential ligand for the ligand binding domain, wherein the novel drug is selected based on its having a greater affinity for the ligand binding domain than that of a known drug.

Figure 1

Statistics from the crystallographic analysis

TABLE 1.

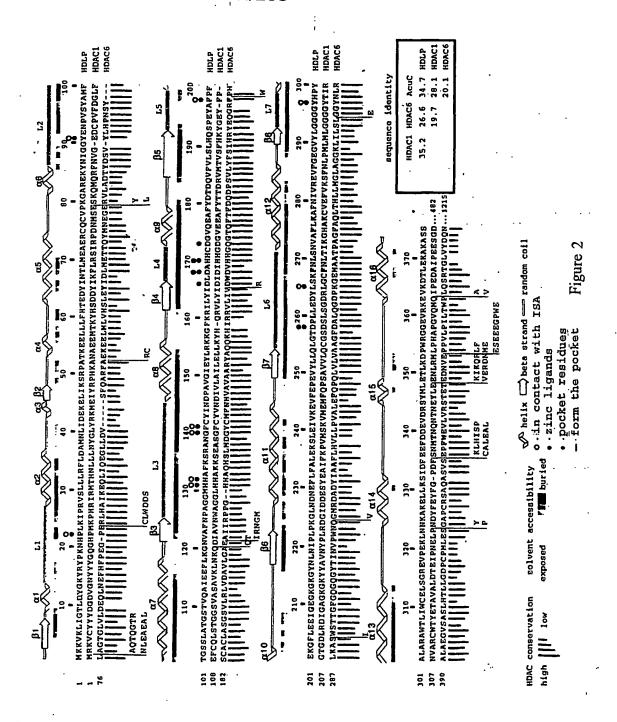
1/263

3-factor 3.55 1.04 180,427 50,796 93.8 RMSD angles .63 .48 125,769 spuoc 0.010 0.009 0.008 Z C2 2.8 27,722 8,753 R-free Aucn 24.0 25.8 25.8 % R-factor C2 3.5 11,454 4,040 86.4 9.6 9.8 22.0 8 Water atoms thimerosal 228 434 456 C2 2.3 79,023 15,958 95.7 atoms Total 3214 3424 6475 Native C2 1.8 134,952 32,143 92.3 2.9 Reflections (IFI > 1a) 31,550 23,582 44,122 WIR analysis (20.0-2.5 Å): Resolution Refinement statistics: د 8 2.0 **(**Y HDLP-Zn-TSA 2.1. Unique reflections Jata coverage (%) phasing power Resolution (Å) Observations Space Group Rcullis (ano) Data set HDLP-Zn Psym (%) **Acullis** Data HDLP Set

Asym = $\Sigma_h \Sigma_l \, ll_h, l - < l_h | > / \Sigma_h \Sigma_l \, l_h, l$ for the intensity (I) of l observations of reflection h. Phasing power = $< F_{\lambda l} > / E$, F_{calc}l/ΣlF_{obs}l, where F_{obs} and F_{calc} are the observed and calculated structure factors, respectively. Figure of merit = IF(hkl)_{best}l/F(hkl). R-free = R-factor calculated using 5% of the reflection data chosen randomly and omitted from the start of refinement. RMSD: root mean square deviations from ideal geometry and root where <F $_{\lambda l}>$ is the root-mean-square heavy atom structure factor and E is the residual lack of closure error. Rcullis is the mean residual lack of closure error divided by the dispersive difference. R-factor = Σ IF $_{
m obs}$ mean square variation in the B-factor of bonded atoms.

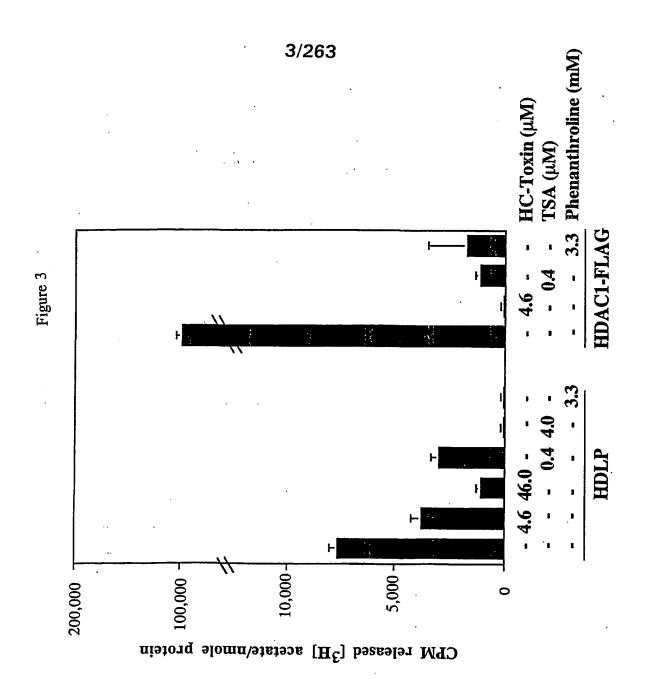
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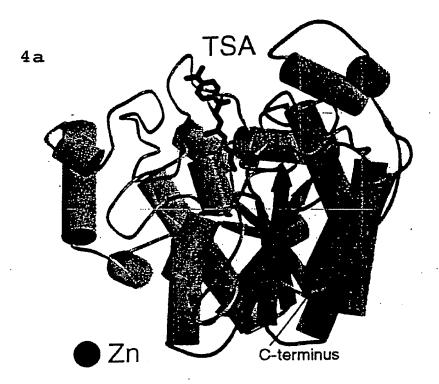
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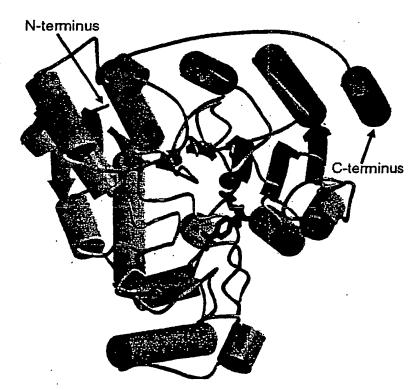
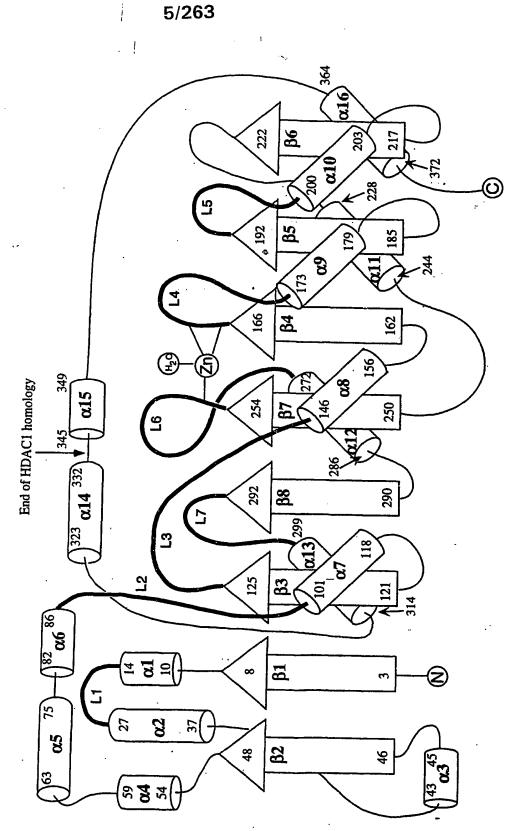


Figure 4
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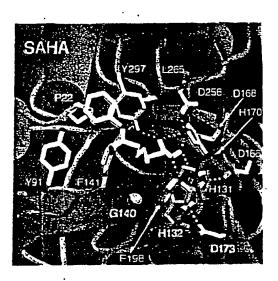
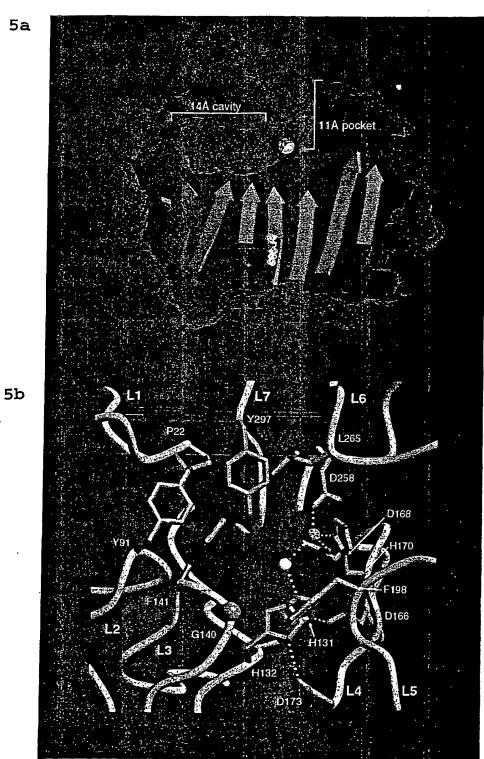


Figure 4D

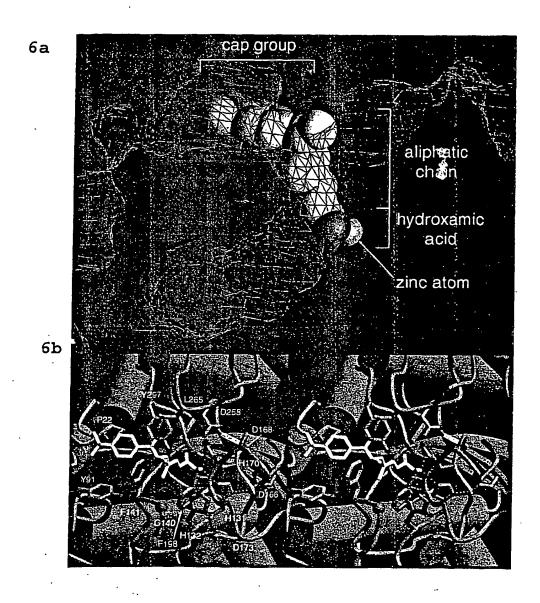
Figure 5 7/263



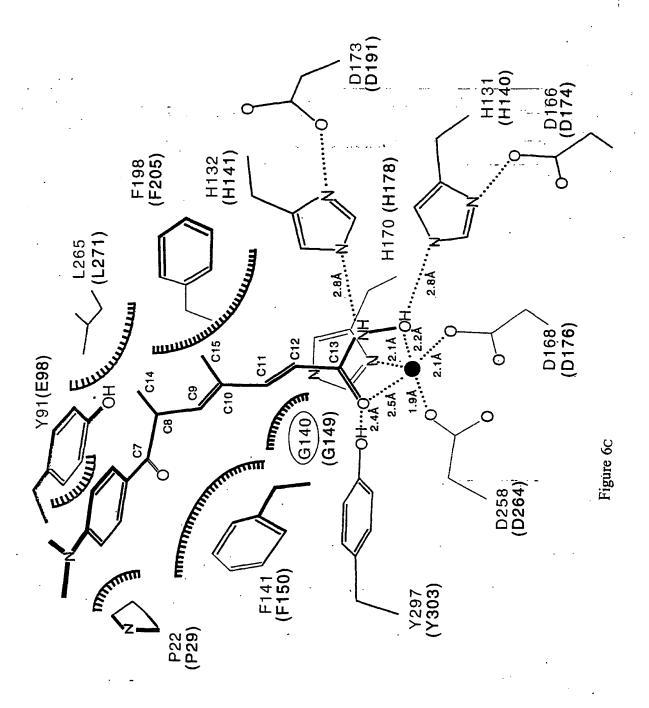
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Figure 6



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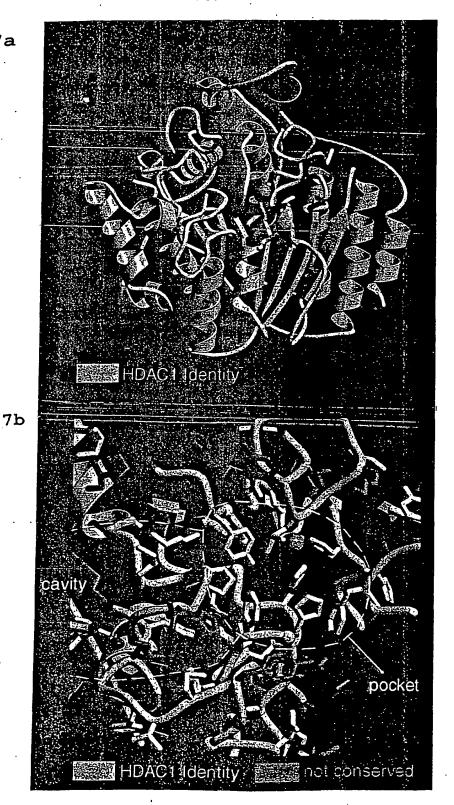
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Figure 7

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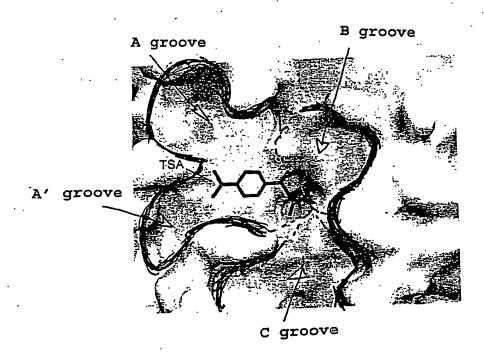


Figure 9

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Figure 10

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Figure 11

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l l	• •
MKKVKLIGILDYGKYRYPKNHPLKIPRVSLLLRFIDAMNI	40
IDEKELIKSRPATKEELLLFHTEDYINTIMEAERCQCVPK	80
GAREKYNIGGYENPVSYAMFTGSSLATGSIVQATEEFLKG	120
NVAFNPAGGMHAFKSRANGFCYINDPAVGIEYLRKKGFK	160
NVAFNPAGATHAF KSIKATOT CTI DETT	200
222	
210 220 230 24	ŧU
	- 240
EKGFLEEIGEGKGKGYNLNIPLPKGLNDNEFLFALEKSLI	: 240 - 220
IVKEVFEPEVYLLQLGIDPLLEDYLSKFNLSNVAFLKAFI	1 280
TVREVEGEGVYLGGGGYHPYALARAWILIWCELSGREVPI	£ 320
KLNNKAKELLKSIDFEEFDDEVDRSYMLETLKDPWRGGE	z 360
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Figure 12

10	20	30	40
•	111111111		
ATGAAGAAGGITAA			
AGTACAGATATCCC			
AGTITICCCTACTCC			
ATAGATGAGAAGGA			
AAGAAGAACTCCIT	-	CGAAGACTA	- •
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CACTITAATGGAAG	CGGAAAGGIG	ICAGIGOGIIV	CCGAAG 240
GGAGCTAGGGAAAA			
CCGTATCTTACGCG	ATGTTTACAG	CICTICICIV	OGCAAC 320
GGGTTCAACAGTGC	'AGGCGATAGA(GAATTTTTA	AAGGGA 360
AATGTAGCTTTCAA	ICCCGCGGGA	GIATGCACC	ACGCTT 400
41 0	42 0	430	44 0
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TTAAAAGCAGGGCA	AACGGCTTTTC	CTACATAAA	CGACCC 440
CGCTGTGGGAATTG	AGTACTTGAG	AAAAAAAGGC	ITTAAG 480
AGAATACTCTACAT	'AGACCITGAT(CCCACCACT(GCGACG 520
GIGITCAGGAAGCC	TTTTACGATAC	CAGACCAGGI	FITCGT 560
CCIGICCCIICACC	AGTCGCCCGAC	TACGCCTTI	CCTTT 600
610	620	630	640
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GAGAAGGGCTTCCT	GGAGGAGATAC	GAGAAGGAA	AAGGAA 640
AGGGCTACAACCTG	AACATTCCCCT	rgccaaaggg	CITGAA 680
CGACAACGAGITCC	TCTTTGCCCTZ	ACAAAAATCTY	CIGGAA 720
ATAGTCAAAGAAGT	ATTIGAGCCCC	AGGITTACC.	PICTIC 760
AACTCGGAACTGAC	CCACICCITG	AGATTACCT:	PICCAA 800
810	820	830	840
GITCAACCICICAAA			
ATCGITCGTGAGGIT			
GAGGCGGATTCCATO			
CCTAATCIGGIGCGA			
AAGCTAAACAATAAA			
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ACTTIGAAGAGTITG			
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AGGAAAGAAGTAAAG			
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Figure 13

10 20 30 40	
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MKKVKLIGILDYGKYRYPKNHPLKIPRVSLILRFLDAMNL	40
IDEKET IKSRPATKEET LI FHTEDYINTIMEAERCQCVPK	80
GAREKYNIGGYENPVSYAMFIGSSLATGSIVQAIEEFLKG	120
NVAFNPAGGMHHAFKSRANGFCYINDPAVGIEYLRKKGFK	160
RILYIDLDAHHCDGVQEAFYDIDQVFVLSLHQSPEYAFPF	200
210 220 230 24	0
<u> </u>	
EKGFLEETGEGKGKGYNLNIPLPKGLNDNEFLFALEKSLE	240
IVKEVFEPEVYLLQLGIDPLLEDYLSKFNLSNVAFLKAFN	280
IVREVFGEGVYLGGGGFHPYALARAWILIWCELSGREVPE	320
KLNNKAKELLKSTDFEFFDDEVDRSYMLETLKDPWRGGEV	360
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17/263 Figure 14

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Figure 15

10	20	30	40	
<u>ll</u>		<u> </u>	لسب	
MKKVKLIGTLDYGE				40
IDEKELIKSRPATK				80
GAREKYNIGGYENE	VSYAMFIGSS.	LATGSTVQA	IEEFLKG	120
NVAFNPAGCMHHAI	KSRANGFCYI	NDPAVGIEY.	LRKKGFK	160
RILYIDLDAHHCD				200
.210	220	230	240)
	<u> </u>	سيلس	لبيبيل	
EKGFLEEIGEGKG				240
IVKEVFEPEVYLL				280
IVREVFGEGVYLG	GGYHPYALAR	AWILIWCEL	SGREVPE	320
1410012 00012-0				
KT NNKAKELLKSII	OFFEFDDEVDR	SYMLETLKD	PWRGGEV	360

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	,					•			
				Residue	* X	Y	Z	OCC. B	Segment ID
ATCM	1	CB	ALA	2	45.336	36.880	75.042	1.00 59.90	AAAA
ATOM	2	c	ALA	2	46.410	38.631	73,628	1.00 52.57	
	3		_	2					AAAA
ATOM		0	ALA		45.780	39.595	74.052	1.00 62.46	AAAA '
ATOM	4	N	ΑLЬ	2	47.540	37.826	75.673	1.00 59.52	AAAA
ATOM	5	CA	ΑĽΑ	2	46.568	37.432	74.527	1.00 57.32	AAAA
ATOM	6	N	LYS	3	46.890	38.570	72.389	1.00 39:51	AAAA
ATOM	7	CA	LYS	3	46.687	39.669	71.440	1.00 29.58	AAAA
ATOM	8		LYS	3	47.855		70.459		
		CE		. 3		39.763		1.00 35.03	AAAA -
ATOM	9	CG	LYS		49.217	40.007	71.102	1.00 55.16	AAAA
ATOM	10	CD	LYS	3	50.315	40.000	73.039	1.00 65.28	AAAA
ATOM	11	CE	LYS	3.	51.700	40.163	70.655	1.00 73.41	AAAA
ATOM	12	NZ	175	3	52.791	40.047	69.642	1.00 69.64	AAAA
ATOM	13	C	LYS	3	45.407	39.422	70.642	1.00 23.29	AAAA
				3					
-ATOM	14	0	LYS		_44.984	38.282	70.487	1.00 27.41	-AAAA
ATOM.	15	N	VAL	4	44.814	40.498	70.138	1.00 25.18	AAAA
ATOM	16	CA	VAL	4	43.585	40.418	59.349	1.00 22.20	AAAA
MOTA	17	CB	VAL	4	42.501	41.365	59.887	1.00 31.46	AAAA
ATOM	18	CG1	VAL	4	41.214	41.202	59.066	1.00 26.55	AAAA
ATOM	19		VAL	4	42.244	41.080	71.348	1.00 34.98	AAAA
				4	43.983				
ATOM	20	C	AYL			40.851	67.961	1.00 25.33	AAAA
ATOM	21	0	VAL	4	44.557	41.927	67.778		AAAA
MOTA	22	N	LYS	5	43.654	40.023	66.978	1.00 21.32	AAAA
ATOM	· 23	CA	LYS	5	44.052	40.291	55.607	1.00 20.10	AAAA
ATOM	24	CB	LYS	5	45.047	39.214	55.177	1.00 23.35	AAAA
ATOM	25	ĊĠ	LYS	5	46.301	39.092	55.049	1.00 23.75	AAAA
ATOM	26	c>	LYS	5	47.183	40.334	£5.919	1.00 23.70	AAAA
ATOM	.27	CE	LYS	Š	48.510	40.151	55.669	1.00 24.34	
				Ś					AAAA
ATOM	28	NZ	LYS	5	49.351	41.357	66.585	1.00 22.04	AAAA
MOTA	29	С	LYS	5	42.914	40.294	54.596	1.00 20.27	AAAA
MOTA	30	0	LYS	5	41.949	39.535	54.728		AAAA
MOTA .	31	N	LEU	6	43.071	41.111	63.564	1.00 19.28	AAAA
MOTA	32	CA	LEU	6	42.097	41.156	52.483	1.00 28.68	AAAA
MOTA	33	CB	LEU	6	41.571	42.574	62.291	1.00 23.51	AAAA
ATOM	34	CG	LEU	6	40.373	42.712	51.342	1.00 30.59	AAAA
ATOM	35		LEU	6	40.079	44.192	51.153	1.00 29.90	AAAA
ATOM	36		LEU	5	40.657	42.085	59.995	1.00 38.98	AAAA
				6					
ATOM	37	C	LEU		42.964	40.701	51.237	1.00 13.17	AAAA
MOTA	38		LEU		. 43.911	41.249	50.919	1.00 22.31	AAAA
MOTA	39	N	ILE	7	42.359	39.689	60.538	1.00 19.15	aaaa
MOTA	40	CA	ILE	7	43.045	39.199	59.338	1. 0 0 13.38	aaaa
ATOM	41	CE	ILE	7	42.922	37.674	59.191	1.00 19.05	AAAA :
ATOM	42	CG2	ILE	7	43.930	37.162	53.144	1.00 16.45	AAAA
ATOM	43		ILE	7	43.253	37.007	60.521	1.00 22.81	AAAA
ATOM	14		ILE	7	43.286	35.543	50.450	1.03 34.99	AAAA
	45	c	ILE	7	42.396	39.850	58.125	1.00 17.95	
ATOM									AAAA .
ATOM	46	0	ILE	7	41.188	39.729	57.928	1.00 19.07	AAAA
ATOM	47	N	GLY	8	43.193	40.562	57.330	1.00 17.70	AAAA
ATOM	48	·CA	GLY	9	42.523	41.193	55.148	1.00 13.11	AAAA
ATOM	49	C	GLY	8	43.640	41.857	55.243	1.00 20.91	AAAA
ATOM	50	D	GLY	8	44.849	41.840	55.504	1.00 22.27	AAAA
ATOM	51	N	THR	9	43.134	42.428	54.155	1.00 23.99	AAAA '
ATOM	52	CA	THR	9	43.950	43.141	53.183	1.00 25.95	AAAA
	53	CE	THR	á	44.739	42.195	52.263	1.00 25.30	AAAA
ATOM									
ATOM	54		THR	9	45.321	42.962	51.199	1.00 25.56	AAAA
MOTA	55		THR	9	43.823	41.144	51.657	1.00 25.24	AAAA ,
ATOM	56	C	THR	9	43.025	43.957	52.294	1.00 29.04	AAAA !
ATOM	57	0	THR	9	41.572	43.582	52.082	1.00 23.05	AAAA
ATOM	58	N	LEU	10	43.517	45.079	51.781	1. 0 0 29. 1 9	AAAA
ATCM	59	CA.	LEU	10	42.690	45.896	50.895	1.00 32.55	AAAA
ATCM	50 50	СВ	LEU	10	43.256	47.319	50.761	1.00 28.09	AAAA
	61	CG	LEU	10	43.142	48.256	51.958	1.00 33.00	AAAA
ATOM						48.403	52.347	1.00 26.65	
ATCM	52	CDT	LEU	10	41.580				AAAA
ATCM	63		LEU	10	43.938	47.744	53.126	1.00 41.33	AAAA
ATOM	64	Ç	LEU	10	42.566	45.261	49.512	1.00 32.68	AAAA.
ATCM	55	0	LEU	10	41.736	45.584	:8.702	1.00 26.97	AAAA
ATCM	ร์ ธ์	N	ASP	11	43.377	44.234	<u> 1</u> 9.256	1.00 25.75	AAAA

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Figure 16-2

MOTA	67	CA	ASP	11	43.367	43.541	47.970	1.00 35.74	AAAA
ATOM	68	СВ	ASP	11	44.477		47.922	1.00 37.61	AAAA
MOTA	69				45.858		48.079	1.00 46.75	AAAA
ATOM	70		1 ASP		46.110		47.444	1.00 46.34	AAAA
ATOM	71	QO	2 ASP	11	46.690		43.821	1.00 58.94	AAAA
MOTA	72	С	ASP	11	42.034		47.607	1.00 34.26	AAAA
ATOM	73		ASP	11	41.748		46.420	1.00 31.12	AAAA
MOTA	74		TYR		41.220		48.609	1.00 26.19	AAAA
MOTA	75				39.923	41.963	48.314	1.00 28.45	AAAA
ATOM	76				39.119	41.720	49.601	1.00 29.35	AAAA
MOTA	77				39.648	40.595	50.470	1.00 28.47	AAAA
ATOM	78			12	40.137		51.755	1.00 32.17	AAAA
MOTA	79			12	40.592	39.808	52.572	1.00.30.35	AAAA
ATOM	80			12	39.629	39.276	50.017	1.00 22.97	AAAA
ATOM	81	CE:		12	40.077	38.228	50.822	1.00 19.60	AAAA
ATOM	82	CZ	TYR	12	40.554	38.499	52.096	1.00 21.42	AAAA
ATOM	83	OH	TYR	12	40.964	37.456	52.907	1.00 23.49	AAAA
ATOM	84	С	TYR	12	39.144	42.907	47.390	1.00 26.67	AAAA
ATOM	85	Ō	TYR	12	38.307	42.466	46.593	1.00 30.51	AAAA
ATOM	86	N	GLY	13	39.441		47.492	1.00 30.22	AAAÄ
ATOM	87	CA	GLY	13	38.767	45.203	46.675	1.00 25.13	AAAA
ATOM	88	С	GLY	13	38.911	45.009	45.177	1.00 27.31	AAAA
MOTA	89	0	GLY	13	38.096	45.522	44.415	1.00 29.38	AAAA
MOTA	90	N	LYS	14	39.937	44.269	44.755	1.00 33.56	AAAA
MOTA	91	CA	LYS	14	40.176	44.005	43.337	1.00 39.81	AAAA
MOTA	92	CB	LYS	14	41.680	44.026	43.031	1.00 51.10	AAAA
MOTA	.93	CG	LYS	14	42.292	45.424	42.907	1.00 64.99	AAAA
MOTA	94	CD	LYS	14	41.757	46.218	41.692	1.00 72.74	AAAA
MOTA	95	CE	LYS	14	42.183	45.639	40.336	1.00 67.25	AAAA
ATOM	96	NZ	LYS	14	41.637	44.280	40.045	1.00 70.06	AAAA
MOTA	97	С	LYS	14	39.589	42.688	42.834	1.00 39.98	AAAA
MOTA	98	0	LYS	14	39.746	42.350	41.658	1.00 46.99	AAAA
MOTA	99	N	TYR	15	38.927	41.944	43.717	1.00 32.64	AAAA
MOTA	100	CA	TYR	15	38.318	40.655	43.355	1.00 41.01	AAAA
ATOM	101	CB	TYR	15	38.996	39.512	44.126	1.00 26.48	AAAA
MOTA	102	CG	TYR	15	40.496	39.571	44.033	1.00 34.97	AAAA
MOTA	103		TYR	15	41.289	39.401	45.167	1.00 43.28	AAAA
MOTA	104		TYR	15	42.677	39.548	45.106	1.00 36.05	AAAA
MOTA	105		TYR	15	41.127	39.879	42.827	1.00 40.78	AAAA
ATOM	106		TYR	15	42.508	40.027	42.756	1.00 37.13	AAAA
MOTA	107	CZ	TYR	15	43.275	39.865	43.899	1.00 36.87	AAAA
MOTA	108 109	OH C	TYR TYR	15 15	44.644 36.838	40.044 40.705	43.844 43.714	1.00 35.40 1.00 38.62	AAAA AAAA
ATOM	110	Ö	TYR	15	36.344	39.868	44.468	1.00 38.02	AAAA
ATOM	111	И	ARG	16	36.141	41.703	43.177	1.00 44.85	AAAA
ATOM	112	CA	ARG	16	34.716	41.890	43.431	1.00 45.75	AAAA
MOTA MOTA	113	CB	ARG	16	34.320	43.348	43.187	1.00 54.17	AAAA
ATOM	114	CG	ARG	16	35.170	44.399	43.875	1.00 66.77	AAAA
ATOM	115	CD	ARG	16	34.920	44.506	45.369	1.00 72.39.	AAAA
ATOM	116	NE	ARG	16	35.649	45.646	45.923	1.00 85.39	AAAA
MOTA	117	CZ	ARG	16	35.489	46.906	45.518	1.00 81.94	AAAA
ATOM	118		ARG	16	34.624	47.197	44.554	1.00 80.19	AAAA
MOTA	119		ARG	16	36.205	47.87.8	46.069	1.00 85.46	AAAA
ATOM	120	C	ARG	16	33.915	41.029	42.460	1.00 43.50	AAAA
MOTA	121	ō	ARG	16	34.400	40.667	41.385	1.00 38.62	AAAA
ATOM	122	N	TYR	17	32.689	40.692	42.833	1.00 32.68	AAAA
ATOM	123	CA	TYR	17	31.850	39.923	41.930	1.00 37.55	AAAA
ATOM	124	СВ	TYR	17	30.662	39.306	42.672	1.00 41.05	AAAA
MOTA	125	CG	TYR	17	31.040	38.104	43.519	1.00 37.51	AAAA
ATOM	126		TYR	17	32.039	38.194	44.493	1.00 32.59	AAAA
ATOM	127	CE1	TYR	17	32.383	37.095	45.277	1.00 29.32	AAAA
ATOM	128	CD2	TYR	17	30.393	36.875	43.346	1.00 31.46	AAAA
MOTA	129		TYR	17	30.726	35.772	44.122	1.00 28.64	AAAA
ATOM	130	CZ	TYR	17	31.721	35.887	45.088	1.00 27.14	AAAA
MOTA	131	ОН	TYR	17	32.044	34.807	45.881	1.00 21.73	AAAA
ATOM	132	C	TYR	17	31.380	40.871	40.836	1.00 40.97	AAAA

ATCM	133	0	TYR	. 17	31.435	42.097	40.984	1.00 29.58	AAAA
ATOM	134		PRO		30.904		39.722	1.00 41.02	
									AAAA
MOTA	135	CD	PRO		30.760	38.910	39.318	1.00 48.67	AAAA
ATOM	136	CA	PRO	18	30.459	41.197	38.649	1.00 49.35	AAAA
ATCM	137	CB	PRO	18	30.321	40.228	37.481	1.00 59.04	AAAA
ATCM	138	CG	PRO	18	29.756	39:017	38.179	1.00 54.15	AAAA
	139.								
ATOM		-	PRO	18	29.178	41.982	38.864	1.00 54.97	AAA A
MOTA	140	0	PRO	18	28.457	41.823	39.850	1.00 46.85	AAAA
ATOM	141	N	LYS	19	28.961	42.868	37.904	1.00 60.87	AAAA
ATOM	142	CA	LYS	19	27.777	43.696	37.749	1.00 67.78	AAAA
MOTA	143	CB	LYS	19	- 27.155	43.278		1.00 73.26	AAAA
ATOM	144	CG	LYS	19	26.971	41.752	36.414	1.00 77.87	AAAA
MOTA	145	CD	LYS	19	26.276	41.166	35.209	1.00 81.01	AAAA
ATOM	146	CE	LYS	19	26.039	39.680	35.471	1.00 82.45	AAAA
	147	NZ	LYS	19	25.417	38.959	34.331	1.00 83.11	
MOTA									AAAA
MOTA	148	C .	LYS	19	26.688	43.594	38.814	1.00 64.15	AAAA
MOTA	149	0	LYS	19	26.810	44.047	39.949	1.00 65.73	AAAA
ATOM	150	N	ASN	20	25.604	42.986	38.345	1.00 59.78	AAAA
ATOM	151	CA	ASN	20	24.353	42.703	39.025	1.00 59.91	AAAA
	152	CB	ASN	20	23.516	41.844			
MOTA							38.077	1.00 68.08	AAAA
ATOM	153	CG	ASN	20	22.108	42.355	37.907	1.00 78.73	AAAA
MOTA	154	OD1	. ASN	20	21.894	43.498	37.496	1.00 78.67	AAAA
MOTA	155	ND2	ASN	20	21.132	41.505	38.211	1.00 83.22	AAAA
ATOM	156	С	ASN	20	24.474	41.977	40.361	1.00 53.35	AAAA
						42.112			
MOTA	157	0	ASN	20	23.611		41.234	1.00 59.92	AAAA
ATOM	158	N	HIS	21	25.543	41.206	40.511	1.00 44.23	AAAA
ATOM	159	CA	HIS	21	25.768	40.397	41.707	1.00 28.15	AAAA
ATOM	160	CB	HIS	21	27.088	39.639	41.570	1.00 31.84	AAAA
ATOM	161	CG	HIS	21	27.155	38.411	42.418	1.00 34.79	AAAA
							43.752		
ATOM T	162		HIS	21	27.344	38.259		1.00 25.03	AAAA
ATOM	163		HIS	21	26.929	37.148	41.917	1.00 34.81	AAAA
ATOM	164	CE1	HIS	21	26.979	36.269	42.900	1.00 17.01	AAAA
MOTA	165	NE2	HIS	21	27.228	36.917	44.026	1.00 32.31	AAAA
ATOM	166	С	HIS	21	.25.763	-41.135	43.051	1.00 29.37	AAAA
	167	Ö							
ATOM			HIS	21	26.346	42.210	43.186	1.00 28.54	AAAA
ATOM	168	N	PRO	22	25.093	40.565	44.066	1.00 29.14	AAAA
MOTA	169	CD	PRO	22	24.301	39.322	44.061	1.00 31.20	AAAA
ATOM	170	CA	PRO	22	25.034	41.185	45.395	1.00 32.84	AAAA
MOTA	171	CB	PRO	22	24.174	40.192	46.187	1.00 34.98	AAAA
	172	CG	PRO	22	23.257	39.634	45.109	1.00 30.11	AAAA
ATOM									
atom	173	C	PRO	22	26.411	41.415	46.044	1.00 34.37	AAAA
ATOM	174	0	PRO	22	26.554	42.272	46.916	1.00 29.17	AAAA
ATOM	175	N	LEU	23	27.415	40.644	45.629	1.00 29.22	AAAA
ATOM	176	CA	LEU	23	28.765	40.781	46.181	1.00 26.49	AAAA
	177	CB	LEU	23	29.414	39.397	46.332		AAAA
ATOM								1.00 22.30	
ATOM	178	CG	LEU	23	28.703	38.527	47.380	1.00 21.04	AAAA
ATOM	179	CD1	LEU	23	29.307	37.113	47.410	1.00 19.35	AAAA
ATOM	180	CD2	LEU	23	28.850	39.197	48.746	1.00 26.51	AAAA
ATOM	181	С	LEU	23	29.661	41.718	45.361	1.00 25.81	AAAA
		ō	LEU		30.893			1.00 28.45	AAAA
ATOM	182			23		41.693	45.477		
MOTA	183	N	LYS	24	29.018	42.539	44.532	1.00 24.86	AAAA
ATOM	184	CA	LYS	24	29.696	43.552	43.723	1.00 27.35	AAAA
ATOM	185	CB	LYS	24	28.662	44.244	42.830	1.00 28.57	AAAA
	186	CG	LYS	24	29.118		42.171	1.00 52.95	AAAA
ATOM							42.283		
ATOM	187	CD	LYS	24	28.025	46.603		1.00 63.74	AAAA
ATOM	188	CE	LYS	24	26.688	46.138	41.706	1.00 66.09	AAAA
ATOM	189	NZ	LYS	24	25.595	47.137	41.896	1.00 66.00	AAAA
ATOM	190	С	LYS	24	30.332	44.592	44.676	1.00 29.52	AAAA
	191	ō	LYS	24	31.412	45.123	44.420	1.00 30.67	AAAA
ATOM									
ATOM	192	N	ILE	25	29.652	44.879	45.779	1.00 26.90	AAAA
ATOM	193	CA	ILE	25	30.151	45.865	46.738	1.00 25.02	AAAA
ATOM	194	CB	ILE	25	29.105	46.177	47.824	1.00 28.34	AAAA
ATOM	195	CG2		25	27.961	46.951	47.237	1.00 23.84	AAAA
		CG1		25	28.661	44.869	48.495	1.00 30.31	AAAA
ATOM	196								
atom	197	CD1		25	27.718	45.051	49.660	1.00 44.90	AAAA
ATOM	198	С	ILE	25	31.424	45.463	47.483	1.00 32.19	AAAA

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MOTA	199	0	ILE	25	31.736	44.271	47.623	1.00 26.54	AAAA
ATOM	200	N	PRO	26	32.191	46.463	47.956	1.00 30.14	AAAA
	201	CD	PRO	26	31.979	47.907	47.770	1.00 36.38	AAAA
MOTA	202	CA	PRO	26	33.431	46.241	48.707	1.00 30.63	AAAA '
ATOM			PRO	26	34.014	47.652	48.814	1.00 34.29	AAAA
MOTA	203	CB				48.373	47.617	1.00 43.39	AAAA
- MOTA	204	CG	PRO	26	33.397			1.00 25.99	AAAA
MOTA	205	С	PRO	26	32.943	45.727	50.061		AAAA
MOTA	206	0	.PRO	26	31.854	46.110	50.484	1.00 25.51	
MOTA .	207	N	ARG	27 .	33.719	44.880	50.743	1.00 21.98	AAAA
MOTA	208	CA	ARG	27	33.267	44.347	52.035	1.00 26.17	AAAA
ATOM	209	CB	ARG	27	32.641	42.969	51.834	1.00 22.70	AAAA
ATOM	210	CG	ARG	. 27	31.442	43.039	50.890	1.00 26.75	AAAA
MÓTA	211	CD	ARG	27	30.832	.41.672	50.581	1.00 33.22	AAAA
ATOM	212	NE	ARG	27	30.121	41.098	51.716	1.00 28.66	AAAA
	213	CZ	ARG	27	30.582	40.129	52.503	1.00 31.79	AAAA
MOTA			ARG	27	31.778	39.598	52.290	1.00 34.08	- AAAA
MOTA	214			27	29.833	39.688	53.505	1.00 26.16	AAAA
MOTA	215		ARG		34.358	44.297	53.090	1.00 24.10	AAAA
MOTA	216	С	ARG	27		45.074	54.038	1.00 23.50	AAAA
MOTA	217	0	ARG	27	34.326			1.00 21.45	AAAA
MOTA	218	N	VAL	. 28	35.314	43.390	52.960		
MOTA	219	CA	VAL	28	36.385	43.385	53.953	1.00 21.75	AAAA
MOTA	220	CB	VAL	28	37.221	42.101	53.866	1.00 26.55	AAAA
ATOM	221	CG1	VAL	28	38.407	42.177	54.830	1.00 23.84	AAAA
MOTA	222	CG2	VAL	28	36.337	40.906	54.214	1.00 19.20	AAAA
MOTA	223	С	VAL	28	37.277	44.611	53.736	1.00 20.86	AAAA
ATOM	224	0	VAL	28	37.770	45.223	54.702	1.00 25.15	AAAA
ATOM	225	N·	SER	29	37.480	44.996	52:475	1.00 19.22	AAAA
MOTA	226	CA	SER	. 29	38.320	46.169	52.209	1.00 19.63	AAAA
	227	CB	SER	29	38.591	46.352	50.702	1.00 24.45	AAAA
MOTA	228	OG	SER	29	37.411	46.697	49.984	1.00 28.74	AAAA
MOTA			SER	29	37.579	47.381	52.756	1.00 21.50	. AAAA
ATOM	229	C		29	38.184	48.320	53.271	1.00 18.95	AAAA
MOTA	230	0	SER		36.256	47.353	52.673	1.00 19.56	AAAA
ATOM	231	N	LEU	30			53.177	1.00 25.97	AAAA
MOTA	232	CA	LEU	30	35.499	48.481	52.744	1.00 22.90	AAAA
MOTA	233	CB	LEU	30	34.032	48.396	53.157	1.00 26.62	AAAA
ATOM	234	CG	LEU	30	33.085	49.541		1.00 28.02	AAAA
MOTA	235		LEU	30	32.885	49.539	54.648		AAAA
MOTA	236	CD2	LEU	30	33.653	50.885	52.698	1.00 25.71	
MOTA	237	C	LEU	30	35.604	48.509	54.696	1.00 18.44	AAAA
ATOM .	238	0	LEU	30	35.704	49.580	55.273	1.00 25.05	AAAA
ATOM	239	N	LEU	31	35.578	47.336	55.336	1.00 19.65	AAAA
ATOM	240	CA	LEU	31	35.672	47.270	56.797	1.00 20.47	AAAA
ATOM	241	CB	LEU	31	35.613	45.821	57.300	1.00 20.60	AAAA
ATOM	242	CG	LEU	31	34.988	45.456	58.665	1.00 39.80	AAAA
MOTA	243		LEU	31	35.712	44.219	59.257	1.00 23.99	AAAA
ATOM	244		LEU	31	35.085	46.591	59.637	1.00 28.48	AAAA
	245	C	LEU	31	37.009	47.870	57.229	1.00 23.85	AAAA
MOTA	246	õ	LEU	31	37.070	48.673	58.154	1.00 21.24	AAAA
ATOM		N	LEU	32	38.079	47.462	56.562	1.00 23.91	AAAA
ATOM	247			32	39.400	47.965	56.899	1.00 24.82	AAAA
ATOM	248	CA	LEU			47.320	56.018	1.00 24.81	AAAA
MOTA	249	CB	LEU	32	40.479		56.276	1.00 27.00	AAAA
ATOM	250	CG		32	40.849	45.854		1.00 27.13	AAAA
MOTA	251		LEU	32	41.995	45.435	55.354	1.00 27.13	AAAA
ATOM	252	CD2	LEU	32	41.285	45.687	57.720		
ATOM	253	С	LEU	32	39.466	49.475	56:763	1.00 19.56	AAAA
MOTA	254	0	LEU	32	39.958	50.143	57.662	1.00 20.71	AAAA
MOTA	255	N	ARG	33	38.974	50.006	55.645	1.00 23.25	AAAA
ATOM	256	CA	ARG	33	39.007	51.449	55.441	1.00 24.33	AAAA
MOTA	257	CB	ARG	33	38.575	51.806	54.013	1.00 23.46	AAAA
	258	CG	ARG	33	39.571	51.327	52.945	1.00 26.94	AAAA
MOTA	259	CD	ARG	33	39.337	51.976	51.585	1.00 42.13	AAAA
MOTA	260	NE	ARG	33	38.023	51.661	51.037	1.00 59.06	AAAA .
MOTA				33	37.583	52.088	49.857	1.00 60.87	AAAA
MOTA	261	CZ	ARG		38.353	52.850	49.095	1.00 65.33	AAAA
MOTA	262	NH1		33			49.433	1.00 56.24	AAAA
MOTA	· 263	NH2		33	36.373	51.743	56.455	1.00 30.33	AAAA
ATOM	264	С	ARG	33	38.124	52.156	20.423	1.00 30.00	,

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ATOM	265	0	ARG	33	38.441	53.252	56,905	1.00 25.45	AAAA
	266		PHE		37.022			1.00 24.98	
ATOM	_								AAAA
MOTA	267		PHE		36.099	52.085		1.00 27.09	AAAA
ATOM	268	CB	PHE	34	34.798	51.276	57.807	1.00 24.88	AAAA
MOTA	269	CG	PHE	34	33.719	51.898	58.631	1.00 20.46	AAAA
ATOM	270		1 PHE	34	33.043	53.018		1.00 18.74	
									AAAA
MOTA	271	CD:		34	33.396	51.383		1.00 20.19	AAAA
ATOM	272	CE	1 PHE	34	32.043	53.627	58.956	1.00 23.04	AAAA
ATOM	273	CE	2 PHE	· 34.	32.406	51.974	60.681	1.00 25.08	AAAA
ATOM	274	CZ	PHE	34	31.726	53.104		1.00 23.31	
									AAAA
ATOM	275	С	PHE	34	36.709	52.115		1.00 23:93	AAAA
ATOM	276	0	PHE	34	36.668	53.138	59.883	1:00 21.71	'AAAA
ATOM	277	N	LYS	35	37.298	51.013	59.645	1.00 21.33	AAAA
ATOM	278	CA	LYS	35	37.862	51.084		1.00 22.54	AAAA
		CB		35		49.716			
MOTA	279		LYS		38.276			1.00 29.70	AAAA
ATOM	280	CG	LYS	35	37.082	48.890		1.00 29.48	AAAA
ATOM	281	CD	LYS	35	37.517	47.535	62.398	1.00 42.17	AAAA
ATOM	282	CE	LYS	35	38.157	46.762	61.275	1.00 34.89	AAAA
ATOM	283	NZ	LYS	35	39.372	47.412	60.719	1.00 67.18	AAAA
ATOM	284	С	LYS	35	39.027	52.055	61.040	1.00 24.68	AAAA
MOTA	285	0	LYS	35	39.282	52.640	62.085	1.00 22.33	AAAA
ATOM	286	N	ASP	36	39.724	52.231	59.926	1.00 25.67	AAAA
ATOM	287	CA	ASP	36	40.842	53.163	59.898	1.00 25.57	AAAA
ATOM	288	CB	ASP	36	41.669	52.984	58.621	1.00 32.26	
									AAAA
MOTA	289	CG	ASP	36	42.881	53.914	58.572	1.00 33.92	AAAA
ATOM	290	ODI	. ASP	36	43.641	53.969	59.563	1.00 40.22	AAAA
ATOM	291	OD2	ASP	36	43.078	54.575	57.538	1.00 40.06	AAAA
ATOM	292	С	ASP	36	40.285	54.578	59.973	1.00 28.04	AAAA
ATOM	293	ō	ASP	36	40.761	55.397	60.765	1.00 29.52	AAAA
ATOM	294.	N	ALA	37	39.272	54.864	59.159	1.00 23.32	AAAA
MOTA	295	CA	ALA	37	38.651	56.192	59.163	1.00 28.22	AAAA
ATOM	296	CB	ALA	37	37.506	56.251	58.119	1.00 25.93	AAAA
ATOM	297	C	ALA	37	38.127	56.549	60.565	1.00 28.41	AAAA
	298	õ	ALA	37	38.186	57.708	60.972	1.00 29.27	AAAA
MOTA									
MOTA	299	N	MET.	38	37.639	55.547	61.300	1.00 24.76	AAAA
ATOM	300	CA	MET	38	37.103	55.727	62.669	1.00 25.45	AAAA
ATOM	301	CB	MET	38	36.077	54.625	62.982	1.00 25.19	AAAA
ATOM	302	CG	MET	38	34.816	54.660	62.148	1.00 22.32	AAAA
ATOM	303	SD	MET	38	33.733	55.983	62.702	1.00 29.90	AAAA
MOTA	304	CE	MET	38	33.402	55.417	64.376	1.00 26.51	AAAA
ATOM	305	С	MET	38	38.203	55.667	63.744	1.00 26.42	AAAA
ATOM	306	0	MET	38	37.924	55.818	64.947	1.00 23.77	AAAA
ATOM	307	N	ASN	39	39.437	55.434	63.300	1.00 26.21	AAAA
ATOM	308	CA	ASN	39	40.607	55.308	64.170	1.00 28.53	AAAA
ATOM	309	CB	ASN	39	40.926	56.643	64.855	1.00 33.95	AAAA
ATOM	310	CG	ASN	39	41.153	57.751	63.858	1.00 29.46	AAAA
ATOM	311	OD1	ASN	39	41.930	57.596	62.925	1.00 36.28	AAAA
ATOM	312	ND2	ASN	39	40.472	58.880	64.046	1.00 40.03	AAAA
ATOM	313	С	ASN	39	40.374	54.223	65.205	1.00 30.07	AAAA
ATOM	314	0	ASN	39	40.682	54.390	66.395	1.00 25.47	AAAA
ATOM	315	N	LEU	40	39.814	53.105	64.744	1.00 28.19	AAAA
ATOM	316	CA	LEU	40	39.527	51.984	65.633	1.00 25.50	AAAA
ATOM	317	CB	LEU	40	38.060	51.562	65.514	1.00 32.14	AAAA
	318	CG	LEU	40	37.044	52.585	66.036	1.00 30.47	AAAA
ATOM									
ATOM	319		LEU	40	35.637	52.027	65.894	1.00 29.07	AAAA
MOTA	320	CD2	LEU	40	37.325	52.889	67.491	1.00 23.80	AAAA
ATOM	321	С	LEU	40	40.433	50.771	65.415	1.00 26.99	AAAA
ATOM	322	0	LEU	40	40.157	49.683	65.915	1.00 25.41	AAAA
							64.691		
MOTA	323	N	ILE	41	41.528	50.970		1.00 28.33	AAAA
ATOM	324	CA	ILE	41	42.459	49.882	64.459	1.00 25.08	AAJA
ATOM	325	CB	ILE	41	42.010	49.020	63.243	1.00 25.01	AAAA
ATOM	326	CG2		41	42.061	49.824	61.961	1.00 22.74	AAAA
	327	CG1		41	42.917	47.802	63.128	1.00 31.01	AAAA
MOTA									
ATOM	328	CD1		41	42.895	46.951	64.341	1.00 42.18	AAAA
ATOM	329	С	ILĒ	41	43.900	50.376	64.247	1.00 24.09	AAAA
ATOM	330	0	ILE	41	44.128	51.406	63.621	1.00 28.92	AAAA
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MOTA	331 N ASI	9 42	44.866	49.634	64.787	1.00 28.95	AAAA
	332 CA ASI		46.279	49.988	64.638	1.00 32.52	AAAA
ATOM					66.007	1.00 34.24	
MOTA				50.094			AAAA
MOTA	334 CG ASI		46.267	51.097	66.911	1.00 51.23	AAAA
MOTA	335 OD1 ASE		46.079	52.250	66.467	1.00 50.19	AAAA
MOTA	336 OD2 ASE		45.924	50.736	68.062	1.00 53.00	AAAA
ATOM	337 C ASI	42	46.985	48.919	63.808	1.00 31.13	. AAAA
ATOM	338 O ASE	42	46.594	47.758	63.838	1.00 26.71	AAAA
ATOM	339 N GLU		48.036	49.312	63.092	1.00 29.99	AAAA
	340 CA GLU		48.793	48.392	62.240	1.00 31.34	AAAA
MOTA							
ATOM	341 CB GLU		50.078	49.045	61.724	1.00 36.12	AAAA
MOTA	342 CG GLU		49.886	50.118	60.676	1.00 52.72	AAAA
MOTA	343 CD GLU		51.214	50.556	60.083	1.00 60.39	AAAA
MOTA	344 OE1 GLU	1 43	51.928	49.688	59.536	1.00 70.32	AAAA
ATOM	345 OE2 GLU	1 43	51.550	51.755	60.163	1.00 60.38	AAAA
ATOM	346 C GLU	. 43	49.196	47.070	62.859	1.00 38.04	AAAA
ATOM	347 O GLU	r 43	49.125	46.024	62.209	1.00 36.83	AAAA
ATOM	348 N LYS		49.636	47.103	64.105	1.00 28.26	AAAA
MOTA	349 CA LYS		50.084	45.879	64.740	1,00 32.71	AAAA
	350 CB LYS		50.974	46.245	65.927	1.00 44.28	AAAA
MOTA						1.00 59.37	
MOTA	351 CG LYS		52.211	47.007	65.418		AAAA
MOTA	352 CD LYS		53.187	47.449	66.491	1.00 68.87	AAAA
MOTA	353 CE LYS		54.373	48.167	65.849	1.00 67.21	AAAA
MOTA	354 NZ LYS	44	55.361	48.648	66.850	1.00 74.00	AAAA
MOTA	355 C LYS	44	48.982	44.889	65.115	1.00 26.75	AAAA
ATOM	356 O LYS	44	49.265	43.792	65.586	1.00 27.37	AAAA
ATOM	357 N GLU	45	47.731	45.278	64.881	1.00 29.20	AAAA
ATOM	358 CA GLU		46.580	44.414	65.165	1.00 21.58	AAAA
ATOM	359 CB GLU		45.387	45.243	65.676	1.00 18.24	AAAA
ATOM	360 CG GLU		45.551	45.828	67.077	1.00 26.57	AAAA
	361 CD GLU		44.418	46.772	67.453	1.00 23.12	AAAA
ATOM			44.224	47.783	66.746	1.00 21.64	AAAA
MOTA	362 OE1 GLU					1.00 26.48	
MOTA	363 OE2 GLU		43.725	46.509	68.454		AAAA
ATOM	364 C GLU			43.710	63.870	1.00 26.31	AAAA
MOTA	365 O GLU		45.400	42.739	63.889	1.00 22.32	AAAA
MOTA	366 N LEU		46.674	44.204	62.748	1.00 20.15	AAAA
MOTA	367 CA LEU		46.317	43.642	61.448	1.00 25.80	AAAA
MOTA	368 CB LEU	46	46.137	44.774	60.433	1.00 27.25	AAAA
ATOM	369 CG LEU	46	45.763	44.397	58.997	1.00 37.72	AAAA
ATOM	370 CD1 LEU	46	44.356	43.810	58.984	1.00 39.46	AAAA
ATOM	371 CD2 LEU	46	45.822	45.632	58.101	1.00 35.43	AAAA
ATOM	372 C LEU	46	47.305	42.623	60.896	1.00 28.88	AAAA
ATOM	373 O LEU	46	48.513	42.860	60.862	1.00 31.98	AAAA
ATOM	374 N ILE	47	46.791	41.469	60.482	1.00 16.92	AAAA
	375 CA ILE	47	47.638	40.448	59.872	1.00 20.98	· AAAA
ATOM			47.412	39.046	67.513	1.00 21.51	AAAA
MOTA	376 CB ILE	47				1.00 20.32	
ATOM	377 CG2 ILE	47	48.115	37.958	52.696		AAAA
MOTA	378 CG1 ILE	47	47.947	39.040	€ .950	1.00 20.71	AAAA
ATOM	379 CD1 ILE	47	49.450	39.207	62.052	1.00 38.87	AAAA
ATOM	380 C ILE	47	47.227	40.417	58.406	1.00 24.50	AAAA
ATOM	381 O ILE	47	46.036	40.279	58.101	1.00 20.74	AAAA
ATOM	382 N LYS	48	48.195	40.550	57.500	1.00 18.73	AAAA
ATOM	383 CA LYS	48	47.883	40.543	56.072	1.00 15.55	AAAA
ATOM	384 CB LYS	48	49.095	40.991	55.239	1.00 16.52	AAAA
MOTA	385 CG LYS	48	48.836	41.011	53.738	1.00 23.25	AAAA ·
	386 CD LYS	48	50.072	41.451	52.957	1.00 32.69	AAAA
ATOM						1.00 26.00	AAAA
ATOM	387 CE LYS	48	49.796	41.496	51.462		
ATOM	388 NZ LYS	48	48.704	42.449	51.114	1.00 46.33	AAAA
MOTA	389 C LYS,	48	47.473	39.140	55.629	1.00 14.43	AAAA
ATOM	390 O LYS	48	48.177	38.174	55.887	1.00 16.83	AAAA
ATCM	391 N SER	49	46.343	39.049	54.945	1.00 16.61	AAAA
ATOM	392 CA SER	49	45.838	37.780	54.439	1.00 14.33	. AAAA
MOTA	393 CB SER	49	44.517	37.984	53.694	1.00 13.21	AAAA
ATOM	394 OG SER	49	43.509	38.613	54.492	1.00 16.86	AAAA
ATOM	395 C SER	49	46.810	37.131	53.459	1.00 24.11	AAAA
	396 O SER	49	47.463	37.815	52.663	1.00 19.59	AAAA
MOTA			2,,	5			

ATOM	1 39	7 N AI	RG 50	ì	46.890	35.805	53.519	1.00 16.83	AAAA
ATOM	1 398	B CA AI	3G 50	1	47.724	35.037	52.610	1.00 23.88	AAAA
ATOM	1 399	9 CB · AI	RG 50	I.	48.805	34.247	53.366	1.00 27.48	
ATOM		CG AF	RG 50		48.284		54.177	1.00 22.99	AAAA
ATOM		L CD AF			49.453		54.759	1.00 25.20	AAAA
ATOM		NE AF			49.073	31.197	55.684	1.00 15.88	AAAA
ATOM					48.411	30.093	55.368	1.00 14.34	AAAA
ATOM					48.023	29.863	54.117	1.00 15.78	AAAA
ATOM					48.150			1.00 16.78	AAAA
ATOM					46.821		51.905	1.00 20.20	AAAA
ATOM	_				45.763	33.650	52.414	1.00 18.63	AAAA
ATOM					47.203	33.596	50.699	1.00 15.63	AAAA
ATOM					48.322	34.028	49.850	1.00 19.45	AAAA
ATOM						32:606	49.994	1.00 14.35	AAAA
ATOM					47.076	32.514	48.629	1.00 17.73	AAAA
ATOM						33.890	48.475	1.00 17.62	AAAA
ATOM	_				46.452	31.256	50.708	1.00 17.02	AAAA
MOTA					47.460		. 51.350	1.00 13.73	AAAA
MOTA					45.377	30.470	50.618	1.00 11.47	AAAA
MOTA					45.375	29.117	51.161	1.00 9.78	AAAA
MOTA					43.967	28.529	51.112	1.00 12.19	
	_				46.301	28.342	50.209	1.00 12.19	AAAA
ATOM					46.307		49.006	1.00 17.19	AAAA
MOTA					47.081	27.392	50.723	1.00 16.40	AAAA AAAA
MOTA					47.952	26.615	49.843	1.00 16.40	AAAA
MOTA					49.109	25.959	50.612	1.00 15.82	AAAA
MOTA					48.582	25.016	51.559	1.00 15.82	AAAA
MOTA MOTA					49.923	27.030	51.336	1.00 16.25	AAAA
ATOM					47.104	25.520	49.215	1.00 14.06	AAAA
ATOM					46.012	25.241	49.690	1.00 14.00	AAAA
ATOM					47.599	24.903	48.145	1.00 17.87	AAAA
ATOM					46.848	23.832	47.492	1.00 19.00	AAAA
MOTA					47.671	23.245	46.339	1.00 13.00	AAAA
MOTA					46.955	22.172	45 539	1.00 22.92	AAAA
ATOM		CD LY			45.787	22.733	44.757	1.00 52.33	AAAA
ATOM		CE LY			46.244	23.565	43.561	1.00 64.17	AAAA
ATOM		NZ LYS			46.898	22.733	42.505	1.00 63.45	AAAA
MOTA		C LY			46.554	22.738	48.520	1.00 22.48	AAAA
ATOM		O LY			45.463	22.158	48.555	1.00 19.97	AAAA
MOTA		N GL			47.536	22.465	49.364	1.00 25.65	AAAA
ATOM		CA GLU			47.389	21.432	50.383	1.00 25.08	AAAA
ATOM		CB GL			48.718	21.241	51.116	1.00 25.40	AAAA
ATOM		CG GLU			48.703	20.185	52.199	1.00 48.95	AAAA
ATOM		CD GLU			50.106	19.821	52.673	1.00 64.21	AAAA
ATOM		OE1 GLU			50.220	19.033	53.640	1.00 62.38	AAAA
ATOM	42	OE2 GLU			51.093	20.311	52.073	1.00 58.22	AAAA
ATOM	.43	C GLU			46.273	21.773	51.362	1.00 18.91	AAAA
ATOM	-44	O GLU			45.489	20.908	51.723	1.00 17.43	AAAA
ATOM	445	N GLU			46.196	23.029	51.786	1.00 16.80	AAAA
MOTA	446	CA GLU			45.137	23.432	52.698	1.00 17.24	AAAA
MOTA	447	CB GLU			45.399	24.855	53.204	1.00 16.15	AAAA
MOTA	448	CG GLU				24.941		1.00 14.41	AAAA
MOTA.	449	CD GLU			47.087	26.354	54.358	1.00 20.17	AAAA
MOTA	450	OE1 GLU		•	46.713	27.252·	53.567	1.00 17.12	AAAA
MOTA	451	OE2 GLU			47.773	26.564	55.394	1.00 18.23	AAAA
	452	C GLU			43.781	23.313	52.000	1.00 15.95	AAAA
ATOM	453	O GLU			42.799	22.869	52.599	1.00 17.82	AAAA
ATOM	454	N LEU			43.722	23.691	50.725	1.00 17.53	AAAA
MOTA	455	CA LEU			42.466	23.579	49.989	1.00 17.33	AAAA
MOTA					42.400	24.177			
ATOM	456 457	CB LEU			42.773	25.707	48.586	1.00 13.86	AAAA
ATOM	457	CG LEU			42.773	26.182	48.552	1.00 15.24 1.00 19.30	AAAA
ATOM	458	CD1 LEU			42.923		47.101		AAAA
ATOM	459	CD2 LEU				26.380	49.207	1.00 15.14	AAAA
ATOM	460	C LEU			42.016	22.126	49.868	1.00 18.46	AAAA
ATOM	461	O LEU	57		40.824	21.823		1.00 17.27	AAAA
ATOM	462	N LEU	58		42.975	21.234	49.636	1.00 16.43	AAAA
		•	-						•

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Figure 16-8

MOTA	463	CA	LEU	58		42.662	19.822	49:475	1.00 15.1	B AAAA
	464	СВ	LEU			43.788	19.113	48.727	1.00 16.0	AAAA •
ATOM			LEU			44.029	19.682	47.321	1.00 21.7	
MOTA	465	CG								
MOTA	466		LEU			45.221	18.982	46.680	1.00 31.9	
MOTA	467	CD2	LEU	58		42.786	19.549	46.469	1.00 34.3	
MOTA	468	С	LEU	58		42.339	19.116	50.787	1.00 21.1	AAAA 6
ATOM	469	0	LEU	58		42.067	17.914	50.795	1.00 20.40	AAAA C
	470	N	LEU			42.377	19.849	51.896	1.00 13.5	
MOTA							19.261	53.173	1.00 15.5	
MOTA	471	CA	LEU			41.958				
ATOM	472		_ LEU	59		42.182	20.236	54.339	1.00 1B.9	
ATOM	473	, CG	PEA	59		43.619	20.537	.54.774	1.00 22.5	7 AAAA
MOTA	474	CD1	LEU	59		43.640	21.654	55.808	1.00 19.8	B AAAA
ATOM	475	CD2	LEU	59	•	44.255	19.253	55.339	1.00 26.7	1 AAAA
	476	C	LEU	59		40.446	18.979	53.043	1.00 17.5	
ATOM	477	ō	LEU	59		39.897	18.112	53.724	1.00 18.0	
ATOM								52.179	1.00 14.6	
MOTA	478	N	PHE	60		39.766	19.737		• • •	
MOTA	479	CA	PHE	60	•	38.338	19.536	51.970	1.00 18.1	
MOTA	480	CB	PHE	60		37.519	20.694	52.557	1.00 18.8	
ATOM	481	CG	PHE	60		36.028	20.564	52.316	1.00 15.9	AAAA 1
ATOM	482	CD1	PHE	60		35.320	19.476	52.817	1.00 19.9	AAAA 8
	483		PHE	60		35.339	21.524	51.576	1.00 18.0	AAAA e
MOTA			PHE			33.947	19.338	52.587	1.00 18.7	
MOTA	484			60						
MOTA	485		PHE	60		33.964	21.399	51.338	1.00 19.1	
ATOM	486	cz	PHE	60		33.268	20.295	51.850	1.00 18.4	
ATOM	487	С	PHE	60		37.916	19.337	50.510	1.00 16.4	
ATOM	488	0	PHE	60		37.227	18.371	50.179	1.00 19.1	AAAA 8
ATOM	489	N	HIS	61	•	38.308	201257	49.638	1.00 18.2	6 AAAA
	490	CA	HIS			37.913	20.163	48.235	1.00 14.4	7 AAAA
ATOM		CB	HIS	61		38.004	21.545	47.582	1.00 17.1	
MOTA	491						22.494	48.084	1.00 14.2	
MOTA	492	CG	HIS	61		36.968				
MOTA	493		HIS	61		35.645	22.580	47.816	1.00 11.0	
ATOM	494	ND1	HIS	61		37.237	23.477	49.012	1.00 23.2	
ATOM	495	CE1	HIS	61		36.121	24.131	49.291	1.00 13.3	
ATOM	496	NE2	HIS	· 61		35.143	23.606	48.579	1.00 21.0	7 AAAA
ATOM	497	C	HIS	61		38.695	19.157	47.417	1.00 18.2	AAAA e
	498	ŏ	HIS	61		39.828	18.819	47.761	1.00 17.5	AAAA 0
ATOM		Ŋ	THR	62		38.071	18.658	46.346	1.00 15.3	
ATOM	499							45.473	1.00 19.0	
MOTA	500	CA	THR	62		38.741	17.686			
MOTA	501	CB	THR	62		37.734	16.767	44.756	1.00 19.6	
ATOM	502	OG1	THR	62		36.795	17.548	44.006	1.00 22.0	
ATOM	503	CG2	THR	62		36.995	15.925	45.767	1.00 28.9	
MOTA	504	С	THR	62		39.595	18.398	44.440	1.00 23.2	2 AAAA
	505	o	THR	62		39.311	19.532	44.044	1.00 17.4	7 AAAA
ATOM	506	N	GLU	63		40.657	17.732	44.009	1.00 18.9	4 AAAA
MOTA						41.571	18.324	43.046	1.00 22.4	
MOTA	507	CA	GLU	63				42.750	1.00 28.3	
ATOM	508	CB	GLU	63		42.736	17.384			
MOTA	509	CG	GLU	63		43.885	17.476	43.708	1.00 60.3	
ATOM	510	CD	GLU	63		45.154	16.893	43.115	1.00 65.0	
ATOM	511	OE1	GLU	63		45.603	17.407	42.065	1.00 66.4	
ATOM	512	OE2	GLU	63		45.697	15.927	43.694	1.00 71.7	2 AAAA
	513	c	GLU	63		40.983	18.764	41.730	1.00 18.6	
MOTA			GLU	63		41.340	19.827	41.228	1.00 18.3	
atom	514	0							1.00 19.7	
ATOM	515	N	ASP	64		40.108	17.943	41.153		
MOTA	516	CA	ASP	64		39.508	18.277	39.864	1.00 .17.8	
ATOM	517	CB	ASP	64		38.584	17.159	39.372	1.00 20.4	
MOTA	518	.CG	ASP	64		37.429	16.884	40.330	1.00 42.7	1 AAAA
	519	OD1		64		36.415	16.291	39.899	1.00 45.0	l AAAA
ATOM	520	OD2		64		37.537	17.243	41.521	1.00 51.7	
ATOM		C		64		38.701	19.582	39.964	1.00 21.9	
ATOM	521		ASP							
ATOM	522	0	ASP	64		38.726	20.410	39.042	1.00 17.3	
ATOM	523	N	TYR	65		37.980	19.750	41.072	1.00 16.1	
ATOM	524	CA	TYR	65		37.178	20.957	41.292	1.00 15.6	
ATOM	525	CB	TYR	65		36.258	20.796	42.529	1.00 12.0	AAAA •
	526	CG	TYR	65		35.501	22.065	42.886	1.00 12.2	
ATOM		CD1		65		34.699	22.718	41.940	1.00 14.7	
ATOM	527							42.253	1.00 18.2	
ATOM	528	CEl	TYR	65		34.028	23.910	94.433	1.00 10.2	- rana

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3 55037	E20	CD3 mVD	65	35.609	22.631	44.163	1.00 13.67	AAAA
MOTA	529	CD2 TYR						
MOTA	530	CE2 TYR	65	34.943	23.824	44.486	1.00 18.16	AAAA
	531	CZ TYR	65.	34.162	24.461	43.533	1.00 16.88	, AAAA
ATOM								
ATOM	532	OH TYR	65	33.555	25.665	43.837	1.00 14.59	AAAA
ATOM	533	C TYR	65	38.090	22.177	41.459	1.00 15.27	AAAA
ATOM	534	O TYR	65	37.882	23.189	40.798	1.00 15.96	AAAA
MOTA	535	N ILE	`66	39.098	22.073	42.321	1.00 14.29	· AAAA
				40.022	23.179	42.540	1.00 18.86	AAAA
MOTA	536	CA ILE	66					
ATOM	537	CB ILE	6 6	41.090	22.836	43.617	1.00 15.56	AAAA
			66	42.152	23.943	43.698	1.00 20.45	AAAA
MOTA	538	CG2 ILE						
MOTA	539	CG1 ILE	66	40.405	22.659	44.967	1.00 19.68	AAAA
	540	CD1 ILE	66	39.717	23.948	45.454	1.00 29.11	· AAAA
MOTA								
MOTA	541	C ILE	66	40,716	23.519	41.236	1.00 25.20	AAAA .
ATOM	542	O ILE	66	40.809	24.692	40.895	1.00 14.60	AAAA
						40.498	1.00 18.21	
MOTA	543	n asn	67	41.190	22.508			AAAA
ATOM	544	CA ASN	6 7	41.879	22.789	39.236	1.00 20.03	AAAA
				42.448	21.523	38.580	1.00 21.73	AAAA
MOTA	545	CB ASN	67					
ATOM	546	CG ASN	67	43.645	20.954	39.333	21.69	AAAA
	547	OD1 ASN	67	44.293	21.645	40.110	1.00 23.97	AAAA
MOTA								
MOTA	548	ND2 ASN	67	43.947	19.692	39.086	1.00 23.23	AAAA
ATOM	549	C ASN	67	40.970	23.500	38.250	1.00 15.87	AAAA
MOTA	550	O ASN	67	41.431	24.347	37.473	1.00 18.64	AAAA
MOTA	551	N THR	68	39.681	23.180	38.295	1.00 16.55	AAAA
						37.400	1.00 20.34	AAAA
MOTA	552	CA THR	68	38.729	23.814			
MOTA	553	CB THR	68	37.360	23.114	37.441	1.00 22.99	AAAA
	554	OG1 THR	68	37.511	21.760	36.978	1.00 21.75	AAAA
ATOM								
ATOM	555	CG2 THR	68	36.378	23.827	36.536	1.00 17.37	AAAA
ATOM	556	C THR	68	· 38.561	25.291	37.755	1.00 16.66	AAAA
	-						1.00 18.79	AAAA
MOTA	557	O THR	68	38.472	26.139	36.871		
ATOM	558	N LEU	69	38.534	25_604	39.045	1.00 14.82	AAAA
			69	38.405	27.000	39.447	1.00 15.20	AAAA
ATOM	559							
ATOM	560	ÇB LEU	69	38.295	27.126	40.973	1.00 16.87	AAAA
ATOM	561	CG LEU	69	37.057	26.551	41.666	1.00 14.76	AAAA
							1.00 16.81	AAAA
MOTA	562	CD1 LEU	69	37.212	26.643	43.179		
ATOM .	563	CD2 LEU	69	35.832	27.312	41.217	1.00 17.26	AAAA
			69	39.623	27.796	38.969	1.00 15.11	AAAA
ATOM	564							
ATOM	565	O LEU	69	39.500	28.934	38.504	1.00 13.30	AAAA
ATOM	566	N MET	70	40.803	27.204	39.090	1.00 13.40	AAAA
						38.659	1.00 16.97	AAAA
MOTA	567	CA MET	70	42.019	27.894			
ATOM	. 568	CB MET	70	43.254	27.114	39.075	1.00 14.87	AAAA
		CG MET	70	43.335	26.886	40.582	1.00 15.18	AAAA
MOTA	569							
ATOM	570	SD MET	70	44.828	25.954	41.060	1.00 28.71	AAAA
ATOM	571	CE MET	70	46.051	27.228	40.893	1.00 21.19	AAAA
						37.155	1.00 19.11	AAAA
ATOM	572	C MET	70	42.064	28.119			
ATOM	573	O MET	70	42.498	29.170	36.700	1.00 17.10	AAAA
	574	N GL	71	41.648	27.118	36.389	1.00 15.06	AAAA
MOTA								AAAA
ATOM	575	CA GLU	71	41.651	27.226	34.934	1.00 16.12	
ATOM	576	CB GLi	71	41.397	25.856	34.305	1.00 16.12	AAAA
				41.387	25.882	32.800	1.00 20.26	AAAA
MOTA	5 7 7	CG GLU	71				and the second s	
MOTA	578	CD GLU	71	42.782	25.920	32.193	1.00 32.31	AAAA
	579	OE1 GLU	71	42.893	25.741	30.958	1.00 27.07	AAAA
ATOM							4 44 44 45	
MOTA	580	OE2 GLU	71	43.762	26.117	32.941	1.00 24.85	AAAA
ATOM	581	C GLU	71	40.580	28.208	34.466	1.00 16.48	AAAA
					29.066	33.611	1.00 17.20	AAAA
MOTA	582	O GLU	71	40.831				
ATOM	583	N ALA	72	39.380	28.097	35.027	1.00 15.68	AAAA
			72	38.300	28.998	34.644	1.00 16.07	AAAA
MOTA	584							
ATOM	585	CB ALA	72	37.035	28.669	35.425	1.00 17.21	AAAA
	586	C ALA	72	38.678	30.453	34.897	1.00 19.07	AAAA
ATOM							1.00 15.92	AAAA
MOTA	587	o ALA	72	38.448	31.326	34.054		
	588	N GLU	73	39.260	30.726	36.062	1.00 15.86	AAAA
ATOM							1.00 15.50	AAAA
MOTA	589	CA GLU	73	39.616	32.097	36.372		
ATOM	590	CB GLU	73	40.046	32.210	37.828	1.00 14.12	AAAA
				40.430	33.615	38.214	1.00 14.24	AAAA
MOTA	591	CG GLU	73					
ATOM ·	592	CD GLU	73	40.961	33.699	39.629	1.00 17.23	AAAA
		OE1 GLU	73	40.147	33.696	40.573	1.00 18.51	AAAA
MOTA	593						1.00 20.88	AAAA
MOTA	594	OE2 GLU	73	42.201	33.753	39.793	1.00 20.00	MANA
•		•						•

MOTA	595	С	GLU	73	40.706	32.709	35.495	1.00 20.36	AAAA
	596	ō	GLU		40.527	33.806	34.948	1.00 17.74	AAAA
MOTA									
MOTA	597	N	ARG		41.832	32.020	35.344	1.00 21.57	AAAA
MOTA	598	CA	ARG	74	42.911	32.623	34.574	1.00 19.48	AAAA
ATOM	599	CB	ARG	74	44.256	31.912	34.834	1.00 18.48	AAAA
		CG	ARG		44.365	30.489	34.351	1.00 14.96	AAAA
MOTA	600								
MOTA	601	CD	ARG	74	45.723	29.892	34.745	1.00 15.05	AAAA
MOTA	602	NE	ARG	74	45.918	28.696	33.950	1.00 18.16	AAAA
ATOM	603	CZ	ARG	· 74	46.439	28.682	32.727	1.00 16.31	AAAA
			LARG	74	46.843	29.811	32.145	1.00 19.74	AAAA
MOTA	604								
MOTA	605	NH2	2 ARG	74	46.466	27.536	32.047	1.00 14.73	AAAA
MOTA	606	С	ARG	74	42.643	32.718	33.084	1.00 16.86	AAAA
ATOM	. 607	0	ARG	74	43.148	33.621	32.426	1.00 15.41	AAAA
	608	N	CYS	75	41.859	31.794	32.547	1.00 17.56	AAAA
ATOM						31.833	31.115	1.00 18.24	AAAA
MOTA	609	CA	CYS	75	41.544				
MOTA	610	CB	CYS	75	41.474	30.414	30.545	1.00 20.94	AAAA
MOTA	611	SG	CYS	75	43.047	29.514	30.572	1.00 19.30	AAAA
ATOM	612	C	CYS	75	40.216	.32.561	30.898	1.00 15.81	AAAA
				75	39.762	32.748	29.762	1.00 17.79	AAAA
MOTA	613	0	CYS						
MOTA	614	N	GLN	76	39.601	32.959	32.007		AAAA
MOTA	615	CA	GLN	76	38.339	33.686	32.010	1.00 23.22	AAAA
A'TOM	616	CB	GLN	76	38.595	35.122	31.530	1.00 22.99	AAAA
				76	37.564	36.107	32.027	1.00 44.69	AAAA
ATOM	617	CG	GLN						
MOTA	618	CD	GLN	76	37.588	36.229	33.535	1.00 47.78	AAAA
MOTA	619	O£1	GLN	76	37.563	35.228	34.243	1.00 62.95	AAAA
MOTA	620	NE2	GLN	76	37.619	37.452	34.033	1.00 45.96	AAAA
	621	C	GLN	76	37.304	32.975	31.135	1.00 23.43	AAAA
MOTA						33.512	30.135	1.00 19.93	· AAAA
MOTA	622	0	GLN	· 76	36.826				
MOTA	623	N	CYS	77	36.951	31.754	31.521	1.00 15.97	AAAA
MOTA	624	CA	CYS	77	36.004	30.979	30.741	1.00 18.91	AAAA
ATOM	625	CB	CYS	. 77	36.738	30.225	29.623	1.00 24.64	AAAA
	626	SG	CYS	77	37.848	28.887	30.269	1.00 25.26	AAAA
MOTA									AAAA
MOTA	627	С	CYS	77	35.302	29.951	31.594	1.00 19.68	
ATOM	628	Ο.	CYS	77	35.685	29.702	32.732	1.00 20.02	AAAA
MOTA	629	N	VAL	78	34.254	29.366	31.022	1.00 16.00	AAAA
MOTA	630	CA	VAL	78	33.531	28.288	31.671	1.00 18.73	AAAA
	631	CB	VAL	78	32.016	28.455	31.557	1.00 15.57	AAAA
MOTA									
MOTA	632		VAL	78	31.312	27.304	32.262	1.00 21.27	AAAA
MOTA	633	CG2	VAL	78	31.603	29.792	32.151	1.00 19.47	AAAA .
ATOM	634	С	VAL	78	33.950	27.077	30.859	1.00 24.02	AAAA
ATOM	635	0	VAL	78	33.499	26.894	29.718	1.00 24.08	AAAA
				79	34.848	26.249	31.420	1.00 18.91	AAAA
MOTA	636	N	PRO						
MOTA	637	CD	PRO	79	35.470	26.341	32.756	1.00 17.70	AAAA
MOTA	638	CA	PRO.	79	35.320	25.056	30.720	1.00 23.37	AAAA
ATOM	639	CB	PRO	79	36.295	24.432	31.732	1.00 21.92	AAAA
	640	CG	PRO	79	36.802	25.677	32.498	1.00 20.90	AAAA
ATOM						24.144	30.376	1.00 27.44	AAAA
MOTA	641	С	PRO	79	34.152				
MOTA:	642	0	PRO	79	33.177	24.064	31.119	1.00 22.20	AAAA
ATOM	643	N	LYS	80	34.245	23.488	29.224	1.00 23.35	AAAA
ATOM	644	CA	LYS	80	33.212	22.570	28.775	1.00 26.78	AAAA
				80	33.708	21.853	27.518	1.00 32.33	AAAA
MOTA	645	CB	LYS			21.055			
ATOM	646	CG	LYS	80	35.098	21.256	27.680	1.00 51.34	AAAA
ATOM	647	CD	LYS	80	35.669	20.817	26.336	1.00 68.70	AAAA
ATOM	648	CE.	LYS	80	37.131	20.401	26.451	1.00 70.04	AAAA
				80	37.688	19.949	25.141	1.00 73.72	AAAA
MOTA	649	NZ	LYS				29.875	1.00 24.71	AAAA
MOTA	650	С	LYS	80	32.875	21.571			
MOTA	65,1	0	LYS	80	33.770	20.957	30.458	1.00 24.23	AAAA
MOTA	652	N	GLY	81	31.582	21.431	30.161	1.00 16.74	AAAA
	653	CA	GLY	81	31.126	20.509	31.194	1.00 18.96	AAAA
MOTA								1.00 22.38	AAAA
ATOM	654	С	GLY	81	31.151	21.039	32.630		
ATOM	655	0	GLY	81	30.604	20.396	33.527	1.00 19.29	AAAA
MOTA	656	N	ALA	82	31.754	22.202	32.863	1.00 22.57	AAAA
	657	CA	ALA	82	31.858	22.738	34.235	1.00 20.65	AAAA
MOTA				82	33.065	23.704	34.333	1.00 20.41	AAAA
ATOM ·	658	CB	ALA						
MOTA	659	C	ALA	82	30.610	23.425	34.781	1.00 21.81	AAAA
MOTA	660	0	ALA	82	30.425	23.529	35.994	1.00 16.95	AAAA '
							-		

MOTA	661	L N	ARG	83	29.758	3 23,926	33.897	1.00 17.6	AAAA 8
MOTA	662				28.549				
ATOM	663				27.777				
	664								
MOTA					26.938				
ATOM	665				26.061				
ATOM	666				25.366)5 AAAA
MOTA	667		-		24.530		36.134	1.00 51.1	.5 AAAA
ATOM	668	NH	II ARG	83	24.286			1.00 55.1	AAAA 0.
MOTA	669	NH	2 ARG	83	23.931	28.646	36.399	1.00 54.2	
ATCM	670	C	ARG	83	27.701	23.530			
MOTA	671	. 0	ARG	83	- 27.193			1.00 24.8	
ATOM	672		GLU	84	27.565			1.00 18.7	
ATOM	673			84	26.768				•
ATOM	674			84	26.527			1.00 24.1	
	675							1.00 32.6	• '
ATOM				84	27.769			1.00 37.9	
ATOM	676				27.832			1.00 51.2	
MOTA	677		1 GLU	84	27.585		30.545	1.00 24.8	2 AAAA
MOTA	678		2 GLU	84	28.114		31.650	1.00 22.5	7 AAAA
ATOM	679	С	GLU	84	27.394		36.043	1.00 25.3	б аааа
MOTA	680	0	GLU	84	26.739	20.321	37.057	1.00 26.1	7 AAAA
ATOM	681	N	LYS	85	28.665	20.232	35.897	1.00 18.7	
ATOM	682	CA	LYS	85	29.399	19.497	36.915	1.00 20.0	
MOTA	683	CB	LYS	85	30.658	18.900	36.280	1.00 18.5	
MOTA	684	CG	LYS	85	31.603	18.223	37.268	1.00 35.6	
ATOM	685	CD	LYS	85	31.151	16.832	37.644	1.00 51.5	
ATOM	686	CE	LYS	85	31.451	15.864	36.520	1.00 59.1	
ATOM	687	NZ	LYS	85	32.914	15.858	36.240	1.00 56.6	
ATOM	688	c	LYS	85	29.811	20.263			
ATOM	689	ō	LYS	85	29.696		38.181	1.00 18.3	
	690	N	TYR	86	30.274	19.738	39.290	1.00 21.6	
ATOM						21.495	38.012	1.00 19.4	
ATOM	691	CA	TYR ·		30.776	22.272	39.145	1.00 14.2	
ATOM	692	CB	TYR	86	32.207	22.692	38.840	1.00 14.9	
ATOM	693	CG	TYR	86	33.107	21.508	38.585	1.00 19.7	б аааа
ATOM	694		TYR	86 .	33.384	20.591	39.601	1.00 18.8	3 AAAA
MOTA	695	CE1		86	34.247	19.519	39.388	1.00 20.2	9 AAAA
ATOM	696	CD2	TYR	86	33.711	21.322	37.337	1.00 18.1	4 AAAA
MOTA	697	CE2	TYR	86	34.567	20.261	37.112	1.00 22.6	AAAA 3
MOTA	698	CZ	TYR	86	34.832	19.364	38.145	1.00 22.6	
ATOM	699	OH	TYR	86	35.680	18.317	37.921	1.00 23.6	
ATOM	700	С	TYR	86	29.967	23.493	39.526	1.00 19.0	
ATOM	701	0	TYR	86	30.353	24.226	40.450	1.00 19.1	
ATOM	702	N	ASN	87	28.873	23.721	38.803	1.00 17.59	
ATOM	703	CA	ASN	87	27.953	24.843	39.071	1.00 18.0	
ATOM	704	CB	ASN	87	27.413	24.730	40.514	1.00 23.8	
ATOM	705	CG	ASN	87	26.020	25.349	40.688		
	706		ASN	87	25.531			1.00 30.6	
ATOM	707		ASN	87		25.520	41.819	1.00 31.55	
ATOM					25.370	25.661	39.580	1.00 20.18	
ATOM	708	C	ASN	· 87	28.641	26.197	38.875	1.00 24.24	
ATOM	709	0	ASN	87	28.283	27.190	39.519	1.00 18.5	
ATOM	710	N	ILE	88	29.617	26.237	37.970	1.00 18.80	
ATCM	711	CA	ILE	88	30.353	27.471	37.680	1.00 18.55	AAAA
ATOM	712	CB	ILE	88	31.865	27.166	37.508	1.00 26.44	AAAA
ATCM	713	CG2	ILE	88	32.613	28.406	37.044	1.00 43.71	L AAAA
ATOM	714	CG1	ILE	88	32.439	26.703·	38.835	1.00 36.30	
ATOM	715	CD1	ILE	88	32.295	27.735	39.888	1.00 24.08	
ATOM	716	С	ILE	88	29.887	28.142	36.392	1.00 14.36	
ATOM	717	ō	ILE	88	29.584	27.459	35.426	1.00 21.93	
	718	N	GLY	89	29.843	29.473			
ATCM	719	CA	GLY	89	29.479		36.380	1.00 18.71	
MOTA						30.162	35.154	1.00 20.23	
atom	720	C	GLY	89	28.147	30.873	35.106	1.00 20.85	
ATCM	721	0	GLY	89	28.006	31.817	34.330	1.00 25.47	
atom	722	N	GLY	90	27.172	30.414	35.889	1.00 21.17	
ATOM	723	CA	GLY	90	25.863	31.060	35.898	1.00 24.44	AAAA
ATOM	724	C	GLY	90	25.862	32.371	36.668	1.00 30.60	AAAA
ATCM	725	0	GLY	90	26.900	32.788	37.168	1.00 28.13	AAAA
ATCM	726	N	TYR	91	24.708	33.036	36.755	1.00 23.38	
							•		

» mon	727 CA TY	R 91	24.598	34.299	37.490	1.00 28.48	3888
ATOM				-			AAAA
MOTA	728 CB TY		23.144	34.753	37.545	1.00 29.88	AAAA
MOTA	729 CG TY	R 91	22.923	35.899	38.518	1.00 33.88	AAAA
MOTA	730 CD1 TY	R 91	23.329	37.197	38.207	1.00 39.69	AAAA
MOTA	731 CE1 TY	R 91	23.130	38.250	39.104	1.00 31.76	AAAA
ATOM-	732 CD2 TY		22.317	35.678	39.759	1.00 40.63	AAAA
	733 CE2 TY		22.115	36.720		1.00 37.07	
ATOM					40.664		AAAA
MOTA	734 CZ TY		22.521	38.002	40.327	1.00 36.22	AAAA
MOTA	735 OH TY	R 91	22.306	39.035	41.210	1.00 44.71	AAAA
MOTA	736 C JY	R 91	25.075	34.157	38.937	1.00 23.59	AAAA
ATOM	737 O TY	R 91	25.713	35.041	39.502	1.00 22.64	AAAA
	738 N GL		24.724	33.032	39.531	1.00 23.09	AAAA
MOTA				32.747			
MOTA			25.048		40.917		AAAA
MOTA	740 CB GL		24.289	31.476	41.306	1.00 32.57	AAAA
ATOM	741 CG GL	U 92	24.595	30.892	42.657	1.00 41.38	AAAA
ATOM	'742 CD GL	J 92	23.604	29.800	43.023	1.00 49.02	- AAAA
ATOM	743 OE1 GL	ງ 92	24.008	28.829	43.715	1.00 45.51	AAAA
	744 OE2 GLT		22.418	29.931	42.628	1.00 38.16	AAAA
MOTA			26.541			-	
MOTA				32.636	41.251	1.00 25.78	AAAA
MOTA	746 O GL		27.045	33.358	42.125	1.00 24.95	AAAA
MOTA	747 N ASI		27.243	31.742	40.556	1.00 21.41	AAAA
ATOM	748 CA ASI	v 93	28.674	31.519	40.777	1.00 21.14	AAAA
ATOM	749 CB ASM	v 93	28.876	30.075	41.226	1.00 17.27	AAAA
MOTA	750 CG ASN		27.905	29.682	42.320	1.00 15.34	AAAA
	751 OD1 ASN		27.882	30.290		1.00 20.33	
ATOM					43.399		AAAA
MOTA	752 ND2 ASN		27.078	28.674	42.047	1.00 20.49	AAAA
MOTA	753 C ASN		29.378	31.778	39.445	1.00 22.25	AAAA
ATOM	754 O ASN	1 93	29.901	30.865	38.806	1.00 20.29	AAAA
ATOM	755 N PRO	94	29.451	33.057	39.045	1.00 25.45	AAAA
ATOM	756 CD PRO	94	29.027	34.221	39.839	1.00 23.03	AAAA
ATOM	757 CA PRO		30.055	33.523	37.794	1.00 23.05	AAAA
			29.669			1.00 28.71	AAAA
MOTA				35.004	37.759		
ATOM	759 CG PRO		28.528		38.755	1.00 40.02	AAAA
ATOM	760 C PRO	94	31.554	33.384	37.697	1.00 26.51	AAAA
MOTA	761 O PRO	94	32.232	33.185	38.688	1.00 17.36	AAAA
ATOM	762 N VAL	95	32.068	33.498	36.478	1.00 21.12	AAAA
ATOM	763 CA VAL		33.506	33.493	36.281	1.00 17.00	AAAA
	764 CB VAL		33.851	33.242	34.796	1.00 25.15	AAAA
MOTA							
ATOM	765 CG1 VAL		35.326	33.537	34.533	1.00 27.19	AAAA
MOTA	766 CG2 VAL		33.551	31.791	34.443	1.00 17.37	AAAA
ATOM	767 C VAL	95	33.989	34.899	36.686	1.00 17.42	AAAA
ATOM	768 O VAL	95	33.426	35.894	36.237	1.00 23.43	AAAA
ATOM	769 N SER	96	34.986	34.982	37.563	1.00 18.84	AAAA
ATOM	770 CA SER		35.564	36.270	37.982	1.00 21.77	AAAA
	771 CB SER		34.608	37.070	38.867	1.00 23.11	AAAA
ATOM							
MOTA	772 OG SER		34.723	36.679	40.223	1.00 24.43	AAAA
ATOM	773 C SER		36.835	35.987	38.789	1.00 29.09	AAAA
ATOM	774 O SER	96	. 37.117	34.828	39.115	1.00 27.12	AAAA
ATOM	775 N TYR	97	37.610	37.020	39.124	1.00 17.51	AAAA
ATOM	776 CA TYR	97	' 38.803	36.751	39.911	1.00 20.69	AAAA
	777 CB TYR	97	39.865	37.835	39.712	1.00 21.82	AAAA
ATOM							
MOTA	778 CG TYR	9.7	40.492	37.748	38.332	1.00 22.72	AAAA
ATOM	779 CD1 TYR	97	39.936	38.414	37.235	1.00 28.47	AAAA
MOTA	780 CE1 TYR	97,	40.473	38.265	35.949	1.00 24.45	AAAA
ATOM	781 CD2 TYR	97	41.599	36.929	38.112	1:00 19.74	AAAA
ATOM	782 CE2 TYR	97	42.144	36.771	36.832	1.00 21.63	AAAA
		97	41.578		35.759	1.00 23.13	AAAA
ATOM	783 CZ TYR			37.439			
ATOM	784 OH TYR	97	42.122	37.273	34.501	1.00 28.54	AAAA
MOTA	785 C TYR	97	38.510	36.515	41.393	1.00 20.12	AAAA
ATOM	786 O TYR	97	39.413	36.285	42.194	1.00 19.76	AAAA
MOTA	787 N ALA	98	37.243	36.558	41.764	1.00 18.56	. AAAA
	788 CA ALA	98	36.899	36.259	43.139	1.00 22.23	AAAA
ATOM		98	35.561	36.888	43.515	1.00 27.92	AAAA
ATOM	789 CB ALA						
· MOTA	790 C ALA	98	36.776	34.743	43.224	1.00 23.56	AAAA
MOTA	791 O ALA	98	36.931.	34.166	44.289	1.00 20.14	AAAA
ATOM	792 N MET	99	36.538	34.094	42.087	1.00 16.84	AAAA
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ATOM	793	CA	MET	99	36.295	32.643	3 42.117	1.00 17.60	2222
	794								AAAA
MOTA		CB	MET		35.864	32.137	7 40.736	1.00 17.05	AAAA
ATOM	795	CG	MET	99	36.999	31.824	1 39.793	1.00 11.16	AAAA
	796	SD	MET	99					
ATOM					36.314	31.698	38.113	1.00 16.54	AAAA
ATOM	797	CE	MET	99	35.165	30.295	38.312	1.00 17.83	AAAA
ATOM	798	С	MET	99	37.432				
							42.650		AAAA
MOTA	799	0	MET	99	37.197	30.753	43.251	1.00 18.21	AAAA
ATOM	. 800	N	PHE	100	38.670				
							42.420	1.00 12.87	AAAA
ATOM	801	CA	PHE	100	39.774	31.439	42.987	1.00 17.13	AAAA
	802	CB	PHE	100					
MOTA					40.559			1.00 15.23	AAAA
ATOM	803	CG	PHE	100	41.647	29.834	42.492	1.00 15.20	AAAA
	804		PHE						
MOTA				100	41.342		43.140	1.00 22.96	AAAA
· ATOM	805	CD2	PHE	100	42.972	30.282	42.488	1.00 17.12	AAAA
	806		PHE	100					
MOTA					42.341	27.901		1.00 19.23	AAAA
ATOM	807	CE2	PHE	100	43.974	29.552	43.129	1.00 16.99	AAAA
	808	CZ				20.200	40		
ATOM			PHE	100	43.658	28.360	43.779	1.00 17.78	AAAA
ATOM	809	С	PHE	100	40.755	32.305	43.774	1.00 20.54	AAAA
	810	o	PHE	100					
MOTA					41.088		44.912	1.00 21.45	AAAA
ATOM	811	N	THR	101	41.219	33.401	43.187	1.00 18.02	AAAA
MOTA	812	CA	THR	101	42.177				
							43.902	1.00 15.25	AAAA
ATOM	813	CB	THR	101	42.715	35.341	42.976	1.00 16.33	AAAA
MOTA	814	OG 1	THR	101	43.386				
								1.00 16.01	AAAA
ATOM	815	CG2	THR	101	43.706	36.226	43.697	1.00 16.31	AAAA
MOTA	816	С	THR	101	41.567	34.860			
								1.00 14.12	AAAA
MOTA	817	0	THR	101	42.110	34.707	46.244	1.00 16.86	AAAA
MOTA	818	N	GLY	102	40.435	35.541	45 000	1.00 13.77	
							45.008		AAAA
MOTA	819	CA	${ t GLY}$	102	39.770	36.145	46.156	1.00 16.29	AAAA
ATOM	820	С	GLY	102	39.330	35.065	47.133	1.00 16.75	
									AAAA
ATOM	821	0	GĹY	102	39.502	35.202	48.338	1.00 14.48	AAAA
ATOM	822	N	SER	103	38.752	33.986	46.615	1.00 16.24	AAAA
ATOM	823	CA	SER	103	38.315	32.890	47.488	1.00 16.72	AAAA
ATOM	824	CB	SER	103	37.567	31.821	46.684	1.00 15.97	AAAA
					27.307				
ATOM	825	OG	SER	103	36.339	32.349	46.197	1.00 26.86	AAAA
ATOM	826	С	SER	103	39.494	32.264	48.218	1.00 17.88	AAAA
	827	0	SER	103					
ATOM					39.405	31.974	49.419	1.00 14.17	AAAA
ATOM	828	N	SER	104	40.604	32.057	47.515	1.00 11.40	AAAA
ATOM	829	CA	SER	104	41.780				
						31.484	48.181	1.00 17.61	AAAA
ATOM	830	CB	SER	104	42.888	31.206	47.160	1.00 15.89	AAAA
ATOM	831	OG	SER	104	42.525	30.102			
							46.362	1.00 27.82	AAAA
ATOM	832	C	SER	104	42.332	32.404	49.271	1.00 17.02	AAAA
MOTA	833	0	SER	104	42.867	31.958	50.286	1.00 15.37	
									AAAA
ATOM	834	N	LEU	105	42.206	33.698	49.052	1.00 17.10	AAAA
ATOM	835	CA	LEU	105	42.709	34.652	50.016	1.00 16.95	AAAA
ATOM		CB ·		105	42.728	36.037	49.365	1.00 18.44	AAAA
ATOM	837	CG	LEU	105	43.613	37.108	49.981	1.00 29.88	AAAA
					45.086				
ATOM		CD1		105	45.080	36 631	49.959	1.00 20.25	AAAA
ATOM	839	CD2	LEU	105	43.438	38 418	49.175	1.00 29,39	AAAA
			LEU	105	41.837				
ATOM						34 637	51.282	1.00 14.81	AAAA
MOTA	841	0 :	LEU	105	42.334	34.703	52.404	1.00 17.74	AAAA
ATOM	842	N.	ALA	106	40.532			•	
						34.531	51.095	1.00 19.28	AAAA
ATOM	843	CA .	àLA	106	39.601	34.493	52.224	1.00 12.39	AAAA
ATOM	844	CB 2	ALA	106	38.140				
						34.574	51.704	1.00 11.58	AAAA
ATOM	845	C i	ALA	106	39.807	33.210	53.023	1.00 14.79	AAAA
ATOM	846	0 3	ALA	106	39.704	33.203	54.250	1.00 13.58	AAAA
ATOM	847 1	N :	THR	107	40.114	32.128	52.318	1.00 13.67	AAAA '
ATOM			THR	107	40.314	30.819	52.956		
						20.013		1.00 13.21	AAAA
MOTA	849 (CB 7	PHR	107	40.187	29.708	51.902	1.00 14.95	AAAA
ATOM			THR	107	38.868	29.792		1.00 15.72	
							51.334		AAAA
ATOM	851 (CG2 7	PHR	107	40.422	28.311	52.511	1.00 9.51	AAAA
			THR	107	41.649				
ATOM						30.751	53.687	1.00 15.80	AAAA
ATOM	853 0)]	CHR	107	41.734	30.206	54.792	1.00 15.63	AAAA
ATOM	854 N		LY	108	42.696	31.294	53.082		
								1.00 14.08	AAAA
ATOM	855 C	CA G	LY	108	43.968	31.298	53.765	1.00 14.62	AAAA
ATOM ·	856		LY	108					
						32.119	55.041	1.00 20.05	AAAA
MOTA	857 C	, 3	LY	108	44.417	31.813	56.063	1.00 17.53	AAAA
ATCM	858 N		ER	109	42.963	33.158	54.988	1.00 15.26	
ALCH	JJJ 1					-J. 130	J3.J00	1.00 13.20	AAAA
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MOTA	85	9 C.	a sei	R 109	42.72	7 34.020	56.153	1.0	0 14.54	. AAAA
MOTA	86	0, C	B SEI	R 109	41.90				0 15.58	
MOTA	86	1 0	G SEI	R 109	42.62				0 16.97	AAAA
MOTA	86	2 C	SE		42.03					AAAA
ATOM	86		SEI		42.189					AAAA
ATOM	86		THE		41.26				0 17.00	AAAA
ATOM	86				40.608					AAAA
	86							1.0	0 12.89	AAAA
ATOM					39.452				0 14.54	AAAA
ATOM	86		31 THE		38.346		-		0 18.11	AAAA
ATOM.	86		32 THE		39.061			1.0	0 12.91	AAAA
MOTA	869		THR		41.633		58.601	1.0	0 18.44	AAAA
MOTA	870		THR		41.574		59.806	1.0	0 16.30	AAAA
MOTA	87:		VAL		42.584	30.013	57.816	1.0	0 15.20	
ATOM	872	2 CA	VAL	111	43.614	29.180	58.403		0 20.45	AAAA
ATOM	873	3 CE	VAL	111	44.517	28.514			0 20.02	AAAA
ATOM	874	l CG	1 VAL	111	45.652	27.765			0 21.79	AAAA
ATOM	875	G CG	2 VAL	`111	43.697				0 19.07	AAAA
ATOM	876	C	VAL	111	44.456				0 18.21	
ATOM	877	7 0	VAL	111	44.838	29.672				AAAA
ATOM	878	N	GLN	112	44.731	31.302	58.890		0 16.82	AAAA
ATOM	879			112	45.493	32.232	59.719			AAAA
ATOM	880			112	45.751	33.540			0 20.13	AAAA
ATOM	881			112	46.593	33.340	58.970		22.39	AAAA
ATOM	882			112			57.723		21.17	AAAA
MOTA	883		1 GLN	112	46.797	34.651	56.982		24.82	AAAA
ATOM	884		2 GLN		47.772	35.381	57.219		25.62	AAAA
	885		GLN	112	45.866	34.963	56.091		13.16	AAĄA
ATOM	886			112	44.743	32.516	61.012		23.99	AAAA
MOTA			GLN	112	45.340	32.593	62.079		17.94	AAAA
MOTA	887	N	ALA	113	43.431	32.700	60.924		15.60	AAAA
ATOM	888	CA	ALA	113	42.653		62.138	1.00	15.04	AAAA
ATOM	889	CB	ALA	113	41.191	33.138	61.802	1.00	18.65	AAAA
MOTA	890	C	ALA	113	42.807	31.751	63.083	1.00	14.84	AAAA
ATOM	891	. 0	ALA	113	42.941	31.909	64.296	1.00	21.05	AAAA
MOTA	892	N	ILE	114	42.767	30.550	62.534	1.00	16.45	AAAA ·
ATOM	893	CA	ILE	114	42.919	29.383	63.389	1.00	15.38	AAAA
ATOM	894	CB	ILE	114	42.600	28.100	62.637	1.00	15.22	AAAA
MOTA	895	CG2		114	42.888	26.893	63.537	1.00	15.72	AAAA
ATOM	896		ILE	114	41.110	28.112	62.244	1.00	19.28	AAAA
ATOM	897	CD1	ILE	114	40.744	27.038	61.191	1.00	13.43	AAAA
MOTA	. 898	C	ILE	114	44.329	29.318	63.968		18.02	AAAA
ATOM	899	0	ILE	114	44.508	28.998	65.156		20.38	AAAA
ATOM ·	900	N	GLU	115	45.328	29.629	63.144		15.27	AAAA
ATOM	901	CA	GLU	115	46.726	29.625	63.614		21.48	AAAA
ATOM	902	CB	GLU	115	47.690	30.080	62.506		21.76	AAAA
ATOM	903	CG	GLU	115	47.884	29.080	61.386		15.78	AAAA
MO.A	904	CD	GLU	115	48.670	29.648	60.211		20.04	AAAA
A COM	905	OE1	GLU	115	49.051	30.843	60.239		21.48	AAAA
A'1'OM	906	OE2	GLU	115	48.901	28.902	59.241		26.59	AAAA
ATOM	907	С	GLU	115	46.877	30.559	64.814		23.55	
ATOM	908	0	GLU	115	47.509	30.212	65.815		23.03	AAAA
ATOM	909	N	GLU	116	46.295	31.748				AAAA
ATOM	910	CA	GLU	116	46.367	32.735	64.703		22.73	AAAA
ATOM	911	CB	GLU	116			65.774		20.54	AAAA
ATOM	. 912	CG	GLU	116	45.744	34.044	65.320		18.40	AAAA
	913	CD	GLU	116	46.562	34.765	64.279		19.76	AAAA
MOTA					47.985	34.998	64.756		27.24	AAAA
ATOM	914		GLU	116	48.164	35.630	65.815		18.44	AAAA
ATOM	915		GLU	116	48.919	34.543			23.17	AAAA
ATOM	916	C	GLU	116	45.682	32.253	67.034		25.39	AAAA
ATOM	917	0	GLU	116	46.207	32.427	68.137	1.00	22.87	AAAA
MOTA	918	N	PHE	117	44.510	31.647	66.872	1.00	18.78	AAAA
MOTA	919	CA	PHE	117	43.778	31.139	68.019		22.11	AAAA
MOTA	920	CB	PHE	117	42.451	30.530	67.581		23.14	AAAA
MOTA	921	CG	PHE	117	41.603	30.054	68.728		24.06	AAAA
MOTA	922	CD1	PHE	117	40.880	30.961	69.493		19.67	AAAA
MOTA	923	CD2	PHE	117	41.559	28.701	69.066		24.08	AAAA
MOTA	924	CE1		117	40.115	30.531	70.586		23.68	AAAA
			-							C-SULVA

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ATOM		HE 117	40.79	99 28.26	70.15	5 1.00 24.04	2222
MOTA	926 CZ P	HE 117	40.07				AAAA
MOTA		HE 117					AAAA
ATOM			44.58				AAAA
		HE 117	44.61	.3 30.03		1.00 24.40	AAAA
ATOM	929 N L	EU 118	45.23	8 29.19		1.00 21.09	
ATOM	930 CA LI	EU 118	46.02				AAAA
ATOM							AAAA
			46.35			1.00 17.90	AAAA
MOTA		EU 118	45.14	8 26.26	4 56.984	1.00 26.20	AAAA
MOTA	933 CD1 LE	EU 118	45.59	1 25.28			
ATOM	934 CD2 LE		44.52				AAAA
ATOM							AAAA
			47.29			1.00 26.49	AAAA
ATOM	936 O LE	EU 118	47.90	8 27.85	6 69.996	1.00 26.34	
MOTA	937 N LY	S 119	47.67	2 29.84		1 00 20 00	AAAA
ATOM	938 CA LY						AAAA
ATOM			48.83				AAAA
			49.39	2 31.61	6 68.805	1.00 30.15	AAAA
ATOM	940 CG LY	S 119	49.91	5 31.26	7 67.437	- 1.00 35.14	
ATOM	941 CD LY	S 119	50.293				AAAA
ATOM	942 CE LY						AAAA
			50.905				AAAA
ATOM	943 NZ LY	S 119	51.199	33.551	l 64.745	1.00 22.46	AAAA
ATOM	944 C LY	S 119	48.335	31.053	70.932	1.00 35.74	
ATOM	945 O LY	S 119	49.117				AAAA
MOTA	946 N GL						AAAA
			47.018			1.00 25.20	AAAA
MOTA	947 CA GL		46.445	31,605	72.309	1.00 30.18	AAAA
ATOM	948 C GL:	Y 120	45.913	33.007		1.00 31.91	
ATOM	949 O GL:	Y 120	45.540			1.00 31.31	AAAA
ATOM	950 N ASI					1.00 34.76	AAAA
			45.889			1.00 20.56	AAAA
MOTA	951 CA ASI		45.353	34.825	70.681	1.00 25.58	AAAA
ATOM	952 CB ASN	J 121	46.278	35.634	69.785	1.00 29.99	
ATOM	953 CG ASN	J 121	47.641				AAAA
ATOM -	954 OD1 ASN		48.396			1.00 24.43	AAAA
ATOM						1.00 54.63	AAAA
			47.944			1.00 41.69	AAAA
ATOM	956 C ASN		43.941	34.759	70.135	1.00 18.85	AAAA
MOTA	957 O ASN	121	43.421	33.675		1.00 24.77	
ATOM	958 N. VAL		43.310			1.00 24.77	AAAA
ATOM						1.00 19.55	AAAA
			41.936	35.994	69.49 9	1.00 22.90	AAAA
ATOM	960 CB VAL		41.053	36.832	70.449	1.00 31.47	AAAA
MOTA	961 CG1 VAL	122	39.649	37.006	69.851		
ATOM	962 CG2 VAL		40.986			1.00 31.52	AAAA
ATOM	963 C VAL			36.154	71.810	1.00 32.50	AAAA
		122	41.953	36.632	68.130	1.00 16.87	AAAA
ATOM	964 O VAL	122	42.518	37.710	67.938	1.00 24.08	AAAA
ATOM	965 N ALA	123	41.321	35.983	67.159	1.00 18.67	
ATOM	966 CA ALA	123	41.360	36.532			AAAA
ATOM	967 CB ALA	123			65.821	1.00 10.18	AAAA
			42.346	35.743	64.990	1.00 19.04	AAAA
ATOM	968 C ALA	123	40.000	36.551	65.131	1.00 13.72	AAAA
ATOM	969 O ALA	123	39.108	35.761	65.439	1.00 20.78	AAAA
ATOM	970 N PHE	124	39.871	37.457	64.180		
MOTA	971 CA PHE	124	38.649			1.00 12.92	AAAA
ATOM	972 CB PHE			37.610	63.405	1.00 14.67	AAAA
		124	37.904	38.878	63.85€	1.00 14.67	AAAA
MOTA	973 CG PHE	124	36.660	39.209	63.049	1.00 20.28	AAAA
ATOM	974 CD1 PHE	124	35.811	38.209	62.587	1.00 18.56	
ATOM	975 CD2 PHE	124	36.286	40.545	62.843	1.00 10.56	AAAA
	976 CE1 PHE				02.043	1.00 19.53	AAAA
MOTA		124	34.609	38.532	61.937	1.00 18.75	AAAA
MOTA	977 CE2 PHE	124	35.072	40.875	62.193	1.00 20.18	AAAA
ATOM	978 CZ PHE	124	34.242	39.867	61.744	1.00 21.57	
MOTA	979 C PHE	124	39.016	37.712	61.930	1.00 21.37	AAAA
ATOM	980 O PHE	124				1.00 22.60	AAAA
			39.823	38.558	61.535	1.00 19.22	AAAA
ATOM	981 N ASN	125	38.449	36.820	61.126	1.00 19.39	AAAA
ATOM	982 CA ASN	125	38.651	36.858	59.691	1.00 16.80	AAAA
ATOM	983 CB ASN	125	39.122	35.507		1 '00 15 54	
ATOM	984 CG ASN	125			59.150	1.00 15.71	AAAA
			39.063	35.469	57.649	1.00 12.84	AAAA
ATOM	985 OD1 ASN	125	39.216	36.508	57.006	1.00 14.91	AAAA
ATOM	986 ND2 ASN -	125	38.853	34.272	57.065	1.00 16.21	
ATCM	987 C ASN	125	37.315	37.210			AAAA
ATOM	988 O ASN	125			JJ.UJ8	1.00 16.22	AAAA
			36.502	36.330		1.00 15.28	AAAA
atom	989 N PRO	126	37.071	38.502	58.775	1.00 14.84	AAAA
ATOM	990 CD PRO	126	37.908	39.684		1.00 18.10	AAAA
•	•	•				+0.10	with
							-

MOTA	991	. CA	PRO	126	35.811	38.910	58.156	1.00 17.33	AAAA
MOTA	992	: CB	PRO	126	35.912	40.434	58.177	1.00 16.32	AAAA
ATOM	993	CG	PRO	126	37.416	40.655	58.008	1.00 20.95	AAAA
MOTA	994	C	PRO	126	35.549	38.359	56.752	1.00 13.78	AAAA
ATOM	995	0	PRO	126	34.404	38.291	56.322	1.00 17.03	AAAA
ATOM -	996		ALA		36.607		56.042	1.00 14.57	AAAA
MOTA	997	CA	ALA		36.463	37.443	54.691	1.00 17.37	AAAA
MOTA	998		ALA		37.816		53.930	1.00 14.48	AAAA
MOTA	999		ALA	127	35.982		54.702	1.00 19.77	AAAA
ATOM	1000	• 0	ALA	127	35.490	35.500	53.688	1.00 15.62	AAAA
ATOM	1001		GLY	128	36.111	35.339	55.849	1.00 13.54	AAAA
MOTA	1002	CA	GLY	.128	35.725	33.939	55.971	1.00 13.53	AAAA
MOTA	1003	С	GLY	128	34.234	33.679	56.101		AAAA
MOTA	1004	0	GLY	128	33.414	34.585	56.017	1.00 15.65	AAAA
MOTA	1005	N	GLY	129	33.883	32.420	56.314	1.00 13.35	AAAA
ATOM	1006	CA	GLY	129	32.487	32.058	56.446	1.00 16.28	-AAAA
ATOM	1007	C	GLY	129	31.754	31.831	55.130	1.00 15.69	AAAA
MOTA	1008	0	GLY	129	30.543	32.021	55.072	1.00 16.10	AAAA
MOTA	1009	N	MET	130	32.479	31.448	54.079	1.00 15.00	AAAA
MOTA	1010	CA	MET	130	31.879	31.163	52.757	1.00 13.35	AAAA
ATOM	1011	CB	MET	130	32.969	31.215	51.689	1.00 12.20	AAAA
MOTA	1012	CG	MET	130	33.680	32.573	51.731	1.00 17.03	AAAA
MOTA	1013	SD	MET	130 130	34.863	32.877	50.425	1.00 15.41 1.00 46.82	AAAA
MOTA	1014	CE	MET MET	130	33.752 31.296	32.973 29.756	49.073 52.885	1.00 40.62	AAAA
MOTA	1015 1016	C Q	MET	130	31.785	28.789	52.297	1.00 12.49	AAAA AAAA
ATOM ATOM	1017	N	HÌS	131	30.188	29.695	53.617	1.00 16.24	AAAA
MOTA	1018	CA	HIS	131	29.556	28.448	54.014	1.00 13.80	AAAA
ATOM	1019	СВ	HIS	131	28.772	28.694	55.316	1.00 15.91	AAAA
ATOM	1020	CG	HIS	131	27.606	29.625	55.175	1.00 13.08	AAAA
ATOM	1021		HIS	131	26.712	30.063	56.096	1.00 12.46	AAAA
ATOM	1022		HIS	131	27.225	30.190	53.976	1.00 22.48	AAAA
ATOM	1023		HIS	131	26.148	30.936	54.166	1.00 16.56	AAAA
ATOM	1024		HIS	131 -	25.817	30.875	55.442	1.00 23.56	AAAA
MOTA	1025	С	HIS	131	28.673	27.663	53.066	1.00 13.69	AAAA
ATOM	1026	0	HIS	131	28.125	26.658	53.470	1.00 17.21	AAAA
ATOM	1027	N	HIS	132	28.523	28.115	51.830	1.00 14.51	AAAA
MOTA	1028	CA	HIS	132	27.669	27.400	50.887	1.00 20.19	AAAA
ATOM	1029	CB	HIS	132	26.863	28.416	50.054	1.00 17.26	AAAA
ATOM	1030	CG	HIS	132	25.748	29.070	50.810	1.00 16.85	AAAA
MOTA	1031		HIS	132	24.787	28.542	51.604	1.00 13.74	AAAA
MOTA	1032		HIS	132	25.497	30.424	50.756	1.00 24.80	AAAA
MOTA	1033		HIS	132	24.429	30.700	51.486	1.00 12.68	AAAA
ATOM	1034		HIS	132 132	23.980	29.576	52.010	1.00 28.65 1.00 16.89	AAAA
ATOM	1035	ć	HIS HIS	132	28.372 27.731	26.412 25.487	49.946 49.460	1.00 14.58	AAAA AAAA
ATOM	1036 1037	. 57	ALA	133	29.669	26.580	49.689	1.00 14.38	AAAA
MOTA	1037	·N CA	ALA	133	30.338	25.680	48.740	1.00 13.76	AAAA
MOTA	1030	CB	ALA	133	31.738	26.194	48.412	1.00 14.95	AAAA
MOTA MOTA	1040	C	ALA	133	30.418	24.219	49.179	1.00 18.80	AAAA
ATOM	1041	Ö	ALA	133	30.557	23.939	50355	.1.00 16.86	AAAA
MOTA	1042	71	PHE	134	30.306	23.306	48.209	1.00 13.76	AAAA
ATOM	1043	ČA	PHE	134	30.378	21.868	48.451	1.00 19.77	AAAA
ATOM	1044	СВ	PHE	134	29.311	21.132	47.620	1.00 15.59	AAAA
ATOM	1045	CG	PHE	134	27.917	21.525	47.975	1.00 17.22	AAAA
ATOM	1046	CD1		134	27.135	22.259	47.091	1.00 17.88	AAA
MOTA	1047	CD2		134	27.392	21.187	49.222	1.00 21.68	AAAA
MOTA	1048	CE1		134	25.836	22.653	47.445	1.00 23.07	AAAA
MOTA	1049		PHE	134	26.099	21.578	49.585	1.00 17.64	AAAA
MOTA	1050	cz	PHE	134	25.323	22.308	48.696	1.00 19.71	AAAA
ATOM	1051	c	PHE	134	31.763	21.354	48.098	1.00 14.76	AAAA
ATOM	1052	5	PHE	134	32.547	22.049	47.442	1.00 18.05	AAAA .
ATOM	1053	N	LYS	135	32.060	20.124	48.515	1.00 16.37	AAAA
ATOM	1054	CΆ	LYS	135	33.369	19.551	48.269	1.00 16.24	AAAA
ATOM	1055		LYS	135	33.360	18.070	48.699	1.00 21.29	AAAA
ATOM	1056	CG	LYS	135	34.640	17.300	48.400	1.00 30.43	AAAA
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Figure 16-17

ATOM	1057	CD	LYS	135	34.59	7 15.86	7 48.977	1.00 30.26	2222
ATOM	1058	CE	LYS	135	34.862			2.00 30.20	AAAA
ATOM			_						AAAA
					36.304				AAAA
ATOM			LYS	135	33.854	19.687	7 46.836		AAAA
MOTA	1061	0	LYS	135	35.020	20.020			
ATOM	1062	N	SER		32.944				AAAA
									AAAA
MOTA		CA	SER		33.301		3 44.490	1.00 15.26	AAAA
MOTA	1064	CB	SER	136	33.339	18.094	43.940		AAAA
ATOM	1065	OG	SER	136	34.135				
ATOM	1066	c	SER	136					AAAA
					32.345				AAAA
MOTA	1067	0	SER	136	32.162	20.071	42.475	1.00 18.77	AAAA
ATOM	1068	N	ARG	137	31.754	21.401			
ATOM	1069	CA	ARG	137	30.805				AAAA
									AAAA
MOTA	1070	CB	ARG	137	29.481	21.448	43.366	1.00 24.19	AAAA
MOTA	1071	CG	ARG	137	28.290	22.273	42.937		AAAA
ATOM	1072	CD	ARG	137	27.026				
									AAAA
MOTA	1073	NE	ARG	137	26.951	20.493	41.862	1.00 50.95	AAAA
ATOM	1074	CZ	ARG	137	26.392	20.781	40.691	1.00 50.38	AAAA
ATOM	1075	NH1	ARG	137	25.854	21.976		1 00 45 56	
ATOM	1076		ARG						AAAA
				137	26.375	19.876		1.00 55.31	AAAA
MOTA	1077	С	ARG	137	30.537	23.595	44.095	1.00 16.14	AAAA
ATOM	1078	0	ARG	137	30.439	23.711		1.00 16.88	
· ATOM	1079	N	ALA	138	30.395			1.00 10.88	AAAA
						24.621		1.00 18.07	AAAA
ATOM	1080	CA	ALA	138	30.117	25.976	43.735	1.00 21.48	AAAA
MOTA	1081	CB	ALA	138	30.460	27.024	42.631	1.00 16.55	AAAA
MOTA	1082	С	ALA	138	28.642	26.090			
MOTA	1083	ō						1.00 21.04	AAAA
			ALA	138	27.798	25.339	43.641	1.00 18.97	AAAA
ATOM	1084	N	ASN	139	28.321	27.019	45.029	1.00 13.83	AAAA
ATOM	1085	CA	ASN	139	26.952	27.158	45.468	1.00 12.92	
ATOM	1086	CB	ASN	139	26.566	25.899	46 274	1.00 12.32	AAAA
								1.00 13.14	AAAA
ATOM	1087	CG	ASN	139	25.162	25.961	46.832	1.00 20.34	AAAA
ATOM	1088	OD1	ASN	139	24.186	26.068	46.086	1.00 19.76	AAAA
ATOM	1089	ND2	ASN	139	25.048		. 48.157		
ATOM	1090	C	ASN	139				1.00 16.36	AAAA
					26.756	28.409	46.315	1.00 20.92	AAAA
ATOM	1091	0	asn	139	27.603	28.738	47.148	1.00 16.81	AAAA
ATOM	1092	N	GLY	140	25.644	29.105	46.086	1.00 19.30	AAAA
ATOM	1093	CA	GLY	140	25.330	30.295			
ATOM	1094						46.864	1.00 21.34	AAAA
			GLY	140	26.393	31.378	46.888	1.00 20.19	AAAA
ATOM	1095	0	GLY	140	26.653	31.968	47.943	1.00 18.77	AAAA
MOTA	1096	N	PHE	141	26.996	31.649	45.733	1.00 15.52	
ATOM	1097		PHE	141	28.034				AAAA
						32.675	45.600	1.00 20.71	AAAA
ATOM	1098		PHE	141	27.711	33.952	46.388	1.00 20.03	AAAA
ATOM	1099	CG	PHE	141	26.355	34.544	46.127	1.00 28.32	AAAA
MOTA	1100	CD1	PHE	141	25.855	35.526	46.997	1.00 24.25	
ATOM	1101	CD2		141	25.589				AAAA
						34.170	45.029	1.00 30.11	AAAA
ATOM	1102	CE1		141	24.628	36.116	46.775	1.00 25.94	AAAA
ATOM	1103	CE2	PHE	141	24.346	34.766	44.801	1.00 21.6:	AAAA
ATOM	1104	CZ :	PHE	141	23.870	35.741	45.677	1.00 24.4	AAAA
ATOM	1105		PHE	141					
					29.357	32.188	46.158	1.00 14.45	AAAA
ATOM		0 1	PHE	141	30.336	32.914	46.111	1.00 16.39	AAAA
ATOM	1107	N (CYS	142	29.389	30.982	46.716	1.00 16.77	AAAA
ATOM	1108	ÇA (rvc	142	30.629	30.466			
							47.285	1.00 17.71	AAAA
MOTA			CYS	142	30.347	29.845	48.659	1.00 13.95	AAA <u>a</u>
ATOM	1110	SG (CYS	142	29.606	30.985	49.846	1.00 16.63	AAAA
ATOM	1111	C C	CYS	142	31.313	29.421	46.401	1.00 18.09	
			YS	142					AAAA
MOTA					30.647	28.527	45.856	1.00 16.60	AAAA
ATOM			YR	143	32.639	29.539	46.272	1.00 12.50	AAAA
ATOM	1114	CA I	YR	143	33.429	28.603	45.478	1.00 15.32	AAAA
ATOM			YR	143	34.333	29.322	44.473		
								1.00 13.07	AAAA
MOTA			YR	143	33.614	30.338	43.612	1.00 15.80	AAAA
ATOM		CD1 T	YR	143	33.396	31.636	44.071	1.00 15.48	AAAA
MOTA	1118 (CE1 T	YR	143	32.740	32.589	43.270	1.00 11.99	AAAA
ATOM		D2 T		143	33.157				
						29.999	42.336	1.00 14.60	AAAA
MOTA		CE2 T		143		30.935	41.532	1.00 10.74	AAAA
ATOM	1121 (Z T	YR	143	32.301	32.229	42.008	1.00 20.89	AAAA
MOTA			YR	143	31.698	33.177	41.208	1.00 18.87	
AI OF		•			J	-3.11	31.200	T.00 TO.01	AAAA
		-							•

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ATOM	1123	C	TYR	143	34.310	27.723	46.358	1.00 17.35	AAAA
MOTA	1124	0	TYR	143	34.581	26.574		1.00 16.67	AAAA
ATOM	1125		ILE		34.763	28.262	_		
	1126								AAAA
MOTA					35.599	27.500		1.00 14.17	AAAA
MOTA	1127		ILE	144	37.018	28.069	48.440	1.00 14.87	AAAA
MOTA	1128	CG	2 ILE	144	37.864	27.332	49.474	1.00 13.55	AAAA
MOTA	1129		1 ILE	144	37.611	28.027		1.00 16.98	
	1130		1 ILE						AAAA
MOTA				144	39.052	28.537		1.00 17.42	AAAA
MOTA	1131	. C	ILE	144	34.959	27.615	49.788	1.00 17.22	AAAA
ATOM	1132	0	ILE	144	34.606	28.716	50.220	1.00 14.72	AAAA
MOTA	1133	N	ASN	145	34.798	26.486		1.00 13.46	AAAA
ATOM	1134			145	34.170				
						26.493	51.797	1.00 16.09	AAAA
MOTA	1135			145	33.401	25.178		1.00 14.50	AAAA
MOTA	1136	CG	ASN	145	32.428	25.239	53.148	1.00 15:64	AAAA
MOTA	1137	OD	1 ASN	145	32.800	25.587	54.263	1.00 14.97	AAAA
ATOM	1138	ND	2 ASN	145	31.170	24.916		1.00 16.74	AAAA
ATOM	1139		ASN	145	35.266	26.639	52.873		
								1.00 15.04	AAAA
ATOM	1140		ASN	145	35.812	25.637	53.338	1.00 15.72	AAAA
MOTA	1141	N	ASN	146	35.599	27.865	53.282	1.00 12.34	AAAA
ATOM	1142	CA	ASN	146	36.685	28.006	54.262	1.00 15.31	AAAA
MÓTA	1143	CB	ASN	146	37.161	29.464	54.354	1.00 15.81	AAAA
ATOM	1144	CG	ASN	146	36.101	30.396	54.865	1.00 15.25	
									AAAA
ATOM	1145		1 ASN	146	36.113	30.757	56.034	1.00 13.57	AAAA
MOTA	1146	ND:	2 ASN	146	35.156	30.775	53.996	1.00 10.85	AAAA
ATOM	1147	C	ASN	146	36.306	27.400	55,613	1.00 13.04	AAAA
ATOM	1148	0	ASN	146	37.160	26.865	56.314	1.00 14.76	AAAA
ATOM	1149	N	PRO	147	35.025	27.489	56.016	1.00 14.28	AAAA
ATOM	1150	CD	PRO	147	33.817	28.175	55.515		
								1.00 7.62	AAAA
MOTA	1151	CA	PRO	147	34.750	26.843	57.307	1.00 13.51	AAAA
MOTA	1152	CB	PRO	147	33.251	27.058	57.482	1.00 14.44	AAAA
MOTA	1153	CG	PRO	147	33.056	28.436	56.827	1.00 12.32	AAAA
MOTA	1154	С	PRO	147	35.118	25.330	57.278	1.00 18.86	AAAA
ATOM	1155	0	PRO	147	35.678	24.796	58.251	1.00 16.24	AAAA
	1156	И	ALA	148	34.818				
ATOM						24.642	56.171	1.00 15.01	AAAA
MOŢA	1157	CA	ALA	148	35.122	23.200	56.080	1.00 15.58	AAAA
MOTA	1158	CB	ALA	148	34.402	22.561	54.882	1.00 12.93	AAAA
ATOM	1159	С	ALA	148	36.624	22.956	55.984	1.00 14.94	AAAA
ATOM	1160	0	ALA	148	37.138	21.999	56.560	1.00 14.69	AAAA
ATOM	1161	N	VAL	149	37.328	23.817	55.263	1.00 12.49	AAAA
	1162	CA	VAL	149	38.778	23.708			
MOTA							55.163	1.00 15.31	AAAA
MOTA	1163	CB	VAL	149	39.364	24.797	54.243	1.00 14.77	AAAA
MOTA	1164	CG1		149	40.899	24.870	54.369	1.00 14.68	AAAA
MOTA	1165	CG2	VAL	149	38.981	24.501	52.808	1.00 12.50	AAAA
ATOM	1166	С	VAL	149	39.323	23.887	56.572	1.00 20.14	. AAAA
ATOM	1167	0	VAL	149	40.172	23.109	57.028	1.00 17.32	AAAA
	1168	N	GLY	50	38.815	24.899	57.271	1.00 15.45	
MOTA									AAAA
MOTA	1169	CA	GLY	50	39.284	25.168	58.622	1.00 20.96	AAAA
ATOM	1170	С	GLY	-50	39.030	24.053	59.621	1.00 24.16	AAAA
MOTA	1171	0	GLY	150	39.888	23.738	60.458	1.00 19.50	AAAA
ATOM	1172	N	ILE	151	37.842	23.465	59.557	1.00 16.67	AAAA
MOTA	1173	CA	ILE	151	37.490	22.375	60.461	1.00 19.56	AAAA
MOTA	1174	CB	ILE	151	35.992	22.052	60.348	1.00 16.46	AAAA
ATOM	1175		ILE	151	35.667	20.709	61.036	1.00 17.93	AAAA
ATOM	1176	CG1	ILE	151	35.180	23.209	60.959	1.00 12.31	AAAA
MOTA	1177	CDI	ILE	151	33.686	23.123	60.672	1.00 18.71	AAAA
ATOM	1178	C	ILE	151	38.352	21.148	60.164	1.00 22.66	AAAA
	1179	ō	ILE	151	38.796	20.472	61.087		
ATOM								1.00 20.08	AAAA
ATOM	1180	N	GLU	152	38.599	20.861	58.888	1.00 19.71	AAAA
ATOM	· 1181	CA	GLU	152	39.434	19.718	58.533	1.00 13.85	AAAA
ATOM	1182	CB	GLU	152	39.362	19.437	57.033	1.00 20.21	AAAA
ATOM	1183	CG	GLU	152	38.033	18.833	56.624	1.00 22.16	AAAA
	1184	CD	GLU	152	37.838	17.430	57.166	1.00 26.94	AAAA
ATOM									
ATOM	1185	0E1		152	36.720	16.906	57.035	1.00 25.03	AAAA
ATCM .	1186	OE2	GLU	152	38.800	16.846	57.708	1.00 24.95	AAAA
ATOM	1187	С	GLU	152	40.865	20.010	58.942	1.00 16.85	AAAA
ATCM	1188	0	GLU .	152	41.629	19.110	59.289	1.00 19.25	AAAA
							•		•

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MOTA	1189	N	.TYR	153		41.228	21.290	58.931	1.00	14.74	AAAA
ATOM	1190	CA	TYR	153		42.574				17.71	AAAA
MOTA	1191	CE	TYR	153		12.757	23.193	59.179	1.00	13.26	AAAA
MOTA	1192	CG	TYR	· 153	4	14.059	23.727	59.729	1.00	16.36	AAAA
MOTA	1193		1 TYR		4	45.234	23.726	58.967	1.00	18.41	AAAA
MOTA	1194					16.438		59.511		21.03	AAAA
ATOM	1195		2 TYR			4.115		61.028		21.16	AAAA
MOTA	1196					15.288		61.570		19.76	AAAA
MOTA	1197					6.440		60.824		25.97	AAAA
ATOM ATOM	1198 1199					17.571		61.410		23.15	AAAA
ATOM	1200		TYR TYR	153		12.712 13.722		60.828 61.247		20.00	AAAA
MOTA	1201		LEU	154		1.683		61.616		19.61 17.78	AAAA
MOTA	1202			154		1.698	21.239	63.042		17.76	AAAA AAAA
ATOM	1203		LEU	154		0.511		63.744		20.44	AAAA
MOTA	1204	CG	LEU			0.636	23.434	63.942		19.57	AAAA
MOTA	1205	CD	1 LEU	154	3	9.277	24.046	64.309		22.48	AAAA
MOTA	1206	CD:	2 LEU	154	4	1.692	23.709	65.044		20.84	AAAA
MOTA	1207	С	LEU	154	4	1.669	19.715	63.262	1.00	19.69	· AAAA
MOTA	1208	0	LEU	154		2.357	19.191	64.149		22.91	AAAA
ATOM	1209	N	ARG	155		0.878	18.996	62.469		20.88	
MOTA	1210	CA	ARG	155		0.840	17.539	62.622		22.64	AAAA
MOTA MOTA	1211 1212	CB CG	ARG	155 155		9.829	16.905	61.652		25.69	AAAA
ATOM	1213	CD	ARG ARG	155		8.384 7.382	17.394 16.834	61.893 60.892		27.64	AAAA
ATOM	1214	NE	ARG	155		6.931	15.497	61.246		25.67 30.88	AAAA AAAA
MOTA	1215	CZ	ARG	155		6.135	14.753	60.488		36.28	AAAA
ATOM	1216		LARG	155		5.705	15.218	59.318		26.96	AAAA
ATOM	1217	NH2	ARG	155		5.737	13.562	60.923		27.33	AAAA
ATOM	1218	С	ARG	155	4	2.235	16.966	62.390		28.00	AAAA
MOTA	1219	0	ARG ·			2.674	16.070	63.119		28.05	AAAA
MOTA	1220	N	LYS	156		2.949	17.486	61.395		23.53	AAAA
ATOM	1221	CA	LYS	156		4.290	16.977	61.128	1.00		AAAA
ATOM	1222	CB	LYS	156		4.854	17.558	59.824		26.01	AAAA
MOTA MOTA	1223 1224	CD	LYS LYS	156 156		6.213 6.632	16.955 17.308	59.444		29.70	AAAA
ATOM	1225	CE	LYS	156		5.685	16.692	58.035 57.005		28.77 39.79	AAAA AAAA
ATOM	1226	NZ	LYS	156		5.671	15.192	57.058		36.33	AAAA
ATOM	1227	C	LYS	156		5.233	17.260	62.299		26.40	AAAA
ATOM	1228	0	LYS	156		6.188	16.511	62.529	1.00		AAAA
MOTA	1229	N	LYS	157 ·	4	4.960	18.337	63.032	1.00	22.50	AAAA
ATOM	1230	CA	LYS	157		5.757	18.709	64.204	1.00	21.12	AAAA
ATOM	1231	CB	LYS	157		5.535	20.181	64.591		28.95	AAAA
ATOM	1232	CG	LYS	157		5.160	21.215	63.652		25.94	AAAA
ATOM	1233 1234	CD	LYS LYS	157 157		7.669 3.281	21.067 22.099	63.575		35.16	AAAA
ATOM ATOM	1235	NZ	LYS	- 157		9.742	21.869	62.627 62.406		39.24 40.01	AAAA AAAA
ATOM	1236	C	LYS	157		5.421	17.825	65.411		22.98	AAAA
ATCM	1237	ō	LYS	157		.085	17.903	66.444		27.77	AAAA
ATOM	1238	N	GLY	158		.392	16.995	65.284		26.49	AAAA
ATOM	1239	CA	GLY	158	44	1.023	16.106	66.376		24.82	AAAA
ATCM	1240	С	GLY	158	42	.771	16.459	67.161	1.00	33.13	AAAA
MOTA	1241	0	GLY	158		.421	15.775	68.128		27.21	AAAA
MOTA	1242	N	PHE	159		.085	17.529	66.781		27.47	AAAA
MOTA	1243	CA	PHE	159		.866	17.861	67.502		24.15	AAAA
ATOM	1244	CB	PHE	159		.410	19.285	67.186		27.53	AAAA
MOTA	1245	CG	PHE	159		.264	20.343	67.827 67.220		27.26	AAAA
MOTA MOTA	1246 1247		PHE PHE	159 159		.439	20.785 20.842	69.076		28.12 21.10	AAAA AAAA
ATOM ATOM	1248	CE1		159		.264	21.714	67.866		26.24	AAAA
ATOM ATOM	1249	CE2		159		.738	21.768	69.736		26.07	AAAA
ATOM	1250	CZ	PHE	159		.907	22.205	69.135		23.91	AAAA
MOTA	1251	C	PHE	159		.792	16.854	67.120		28.02	AAAA
ATCM	1252	0	PHE	159		.639	16.533	65.947		21.14	AAAA
ATCM	1253	N	LYS	160 '		.056	16.361	68.110	1.00	24.79	AAAA
MOTA	1254	CA	LYS	160	38	.011	15.366	67.881	1.00	24.26	AAAA
•								-			

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MOTA	1255	CB	LYS	160	38.360	14.098	68.668	1.00	22.86	AAAA
MOTA	1256	CG	LYS	160	39.625	13.424	68.157		43.16	AAAA
ATOM	1257	CD	LYS	160	40.222		69.141		54.05	AAAA
MOTA	1258	CE	LYS	160	39.236		69.577		62.87	AAAA
ATOM	1259	NZ	LYS	160	38.154		70.446		68.11	AAAA
ATOM-	1260	С	LYS		36.599		68.225		21.12	AAAA
ATOM	1261		LYS		35.632		68.051		22.43	AAAA
ATOM	1262		ARG		36.476		68.733		19.68	AAAA
ATOM	1263		ARG		35.164		69.073		20.84	AAAA
ATOM	1264		ARG		34.865		70.572		26.02	AAAA
ATOM	1265		ARG		34.715		71.080		28.47	AAAA
ATOM	1266		ARG		34.213		72.523			AAAA
MOTA	1267		ARG		35.098		73.445		32.99	. AAAA
ATOM	1268		ARG		36.272		73.883	_	40.49	AAAA
MOTA	1269		1 ARG		36.724		73.489		31.49	AAAA
MOTA	1270	NH:	2 ARG	161	37.003	17.014	74.712		38.54	- AAAA
ATOM	1271	С	ARG	161	35.171	19.060	68.680		18.98	AAAA
ATOM	1272	0	ARG	161	35.552	19.932	69.460		23.57	AAAA
ATOM	1273	N	ILE	162	34.743	19.332	67.458		19.82	AAAA
MOTA	1274	CA	ILE	162	34.744	20.700	66.947		17.81	AAAA
ATOM	1275	CB	ILE	162	35.522	20.717	65.626		18.33	AAAA
ATOM	1276	CG2	: ILE	162	35.542	22.110	65.042	1.00	13.65	AAAA
ATOM	1277	CG1	ILE	162	36.937	20,200	65.895	1.00	18.15	AAAA
ATOM	1278	CD1	LILE	162	37.722	19.852	64.670	1.00	22.52	AAAA
MOTA	1279	С	ILE	162	33.316	21.184	66.724	1.00	14.71	AAAA
MOTA	1280	0	ILE	162	32.520	20.492	66.126	1.00	17.99	AAAA
MOTA	1281	N	LEU	163	32.996	22.374	67.217	1.00	16.93	AAAA
ATOM	1282	CA	LEU	163	31.653	22.902	67:061	1.00	20,.73.	АЛАА
MOTA	1283	CB	LEU	163	31.115	23.376	68.421		18.45	AAAA
MOTA	1284	CG	LEU	163	29.846	24.236	68.463			AAAA
MOTA	1285		LEU	163	28.657	23.408	67.975		15.66	AAAA
MOTA	1286		LEU	163	29.609	24.751	69.870		18.74	AAAA
MOTA	1287	C	LEU	163	31.705	24.071	66.106		18.40	AAAA
MOTA	1288	0	LEU	163	32.607	24.889	66.188		18.65	AAAA
ATOM	1289	N	TYR	164	30.752	24.128	65.186		16.97	
MOTA	1290	CA	TYR	164	30.656	25.246	64.252		11.76	AAAA
ATOM	1291	CB	TYR	164	30.782	24.754	62.816		14.07	AAAA
MOTA	1292 1293	CG CD1	TYR TYR	164 . 164	30.593 31.573	25.851	61.797		14.51	AAAA
MOTA MOTA	1294	CE1		164	31.353	26.822 27.832	61.562		27.08	AAAA
MOTA	1295	CD2		164	29.415	25.916	60.598 61.070		26.21 21.45	AAAA AAAA
ATOM	1296	CE2		164	29.193	26.891	60.137		21.45	AAAA
ATOM	1297	CZ	TYR	164	30.148	27.839	59.896		16.35	AAAA
ATOM	1298	ОН	TYR	164	29.857	28.764	58.913		27.44	AAAA
ATOM	1299	C.	TYR	164	29.279	25.873	64.463		15.67	
ATOM	1300	ō	TYR	164	28.760	25.177	64.455		16.07	AAAA
ATOM ·	1301	N	ILE	165	2940	27.187	64.674		14.52	AAAA
ATOM	1302	CA	ILE	165	27.178	27.887	64.893		18.37	AAAA
ATOM	1303	CB	ILE	165	27.959	28.596	66.254		13.31	AAAA
ATOM	1304	CG2	ILE	165	26.654	29.359	66.419		13.06	AAAA
ATOM	1305	CG1		165	28.172	27.573	67.376		17.28	AAAA
ATOM	1306	CD1		165	28.493	28.209	68.739		15.02	AAAA
ATOM	1307	С	ILE	165	27.853	28.926	63.779		20.75	AAAA
ATOM	1308	0	ILE	165	28.759	29.733	63.569		1.6.67	AAAA
ATOM	1309	N	ASP	166	26.725	. 28.901	63.084	1.00	15.37	AAAA
ATOM	1310		ASP	166	26.503	29.779	61.942		15.63	AAAA
ATOM	1311	СВ	ASP	166	26.276	28.885	60.698		12.31	AAAA
MOTA	1312		ASP	166	26.279	29.666	59.393	1.00		AAAA
ATOM	1313	OD1		166	25.378	30.508	59.213	1.00		AAAA
ATOM	1314	OD2	ASP	166	27.187	29.428	58.551		16.06	AAAA
MOTA	1315	С	ASP	166	25.334	30.740	62.174	1.00	15.54	AAAA
MOTA	1316	0	ASP	166	24.160	30.355	62.137		12.60	AAAA
MOTA	1317		LEU	167	25.647	32.010	62.407	1.00		AAAA
ATOM	1318		LEU	167	24.598	32993	62.665	1.00	12.05	AAAA
MOTA	1319		LEU	167	25.051	33.962	63.767	1.00	14.61	AAAA
à T CM	1320	CG	LEU	167	25.345	. 33.239	65091	1.00	17.20	AAAA

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MOTA	132:	L C	D1 LEU	167	25.63	5 34.271	66.169	1.00 28.82	AAAA
ATOM	1322	C1	D2 LEU		24.14				
									AAAA
ATOM	1323		LEU		24.12	2 33.776	61.449	1.00 12.62	AAAA
ATOM	1324	1 0	LEU	167	23.28	34.678	61.570	1.00 15.00	AAAA
ATOM	1325	N	ASP	168	24.66				
								1.00 14.35	AAAA
ATOM	1326			168	24.27		59.022	1.00 19.50	AAAA
MOTA	1327	CI CI	3 ASP	168	25.060	33.409	57.880	1.00 25.15	AAAA
ATOM	1328	C	ASP	168	24.908				
			_					1.00 48.45	AAAA
ATOM	1329		O1 ASP	168	25.47		56.454	1.00 64.45	AAAA
ATOM	1330	OI	D2 ASP	168	24.215	33.633	55.668	1.00 44.71	AAAA
ATOM	1331	. с	ASP	168	22.787		58.834	1.00 16.30	
MOTA	1332								AAAA
			ASP	168	22.327		59.252	1.00 17.72	AAAA
MOTA	1333	N	ALA	169	22.059	34.657	58.175	1.00 14.11	AAAA
MOTA	1334	CA	ALA	169	20.618	34.503	57.934	1.00 19.61	AAAA
MOTA	1335			169	20.006				
							57.470	1.00 13.56	AAAA
ATOM	1336	С	ALA	169	20.277	33.400	56.926	1.00 18.23	AAAA
MOTA	1337	0	ALA	169	19.105	33.159	56.641	1.00 17.20	AAAA
ATOM	1338	N	HIS	170	21.301				
							56.3 7 3	1.00 16.53	AAAA
MOTA	1339	CA	HIS	170	21.075	31.652	55.436	1.00 17.51	AAAA
MOTA	1340	CB	HIS	170	21.616	31.973	54.033	1.00 22.32	AAAA
ATOM	1341	CG	HIS	170	20.954		53.377		
								1.00 25.38	AAAA
ATOM	1342		2 HIS	170	19.934		52.487	1.00 19.33	AAAA
MOTA	1343	ND	1 HIS	170	21.308	34.448	53.638	1.00 18.17	AAAA
MOTA	1344	CE	1 HIS	170	20.535	35.257	52.935	1.00 30.34	AAAA
ATOM	1345		2 HIS	170	19.692				
							52.229	1.00 17.51	AAAA
ATOM	1346	C	HIS	170	21.781		55.967	1.00 16.72	AAAA
ATOM	1347	0	HIS	170	22.827	30.511	56.610	1.00 15.92	AAAA
ATOM	1348	N	HIS	171	21.209	29.245	55.682	1.00 15.28	
ATOM	1349	CA.							ÀAAA
				171	21.751	27.961	56.123	1.00 12.53	AAAA
ATOM	1350	CB	HIS	171	20.702	26.878	55.814	1.00 14.09	AAAA
ATOM	1351	CG	HIS	171	21.180	25.468	55.980	1.00 17.27	AAAA
ATOM	1352	CD.	2 HIS	171	21.249	24.447			
							55.090	1.00 12.48	AAAA
ATOM	1353		l HIS	171	21.622	24.956	57.181	1.00 26.73	AAAA
MOTA	1354	CE:	L HIS	171	· 21.948	23.685	57.021	1.00 15.98	AAAA
ATOM	1355	NE:	2 HIS	171	21.729	23.352	55.761	1.00 20.03	AAAA
ATOM	1356	C	HIS	171	23.107			•	
						27.602	55.498	1.00 15.55	AAAA
MOTA	1357	0	HIS	171	23.318	27.784	54.298	1.00 17.03	AAAA
ATOM	1358	N	CYS	172	24.026	27.105	56.323	1.00 14.33	AAAA
ATOM	1359	CA	CYS	172	25.350	26.675	55.866	1.00 13.65	AAAA
ATOM	1360	CB	CYS	172	26,330				
						26.631	57.054	1.00 12.99	AAAA
ATOM	1361	SG	CYS	172	25.680	25.826	58.551	1.00 17.17	AAAA
MOTA	1362	С	CYS	172	25.212	25.274	55.257	1.00 16.52	AAAA
ATOM	1363	0	CYS	172	25.750	24.297	55.783	1.00 14.95	AAAA
ATOM	1364	N	ASP	173					
					24.516	25.173	54.130	1.00 15.42	AAAA
ATOM	1365	CA	ASP	173	24.302	23.865	53.531	1.00 14.75	'AAAA
MOTA	1366	CB	ASP	173	23.339	23.956	52.332	1.00 17.73	AAAA
MOTA	1367	CG	ASP	173	23.765	24.966	51.283	1.00 22.84	AAAA
ATOM	1368		ASP	173	23.106				
						24.998	50.216	1.00 18.68	AAAA
ATOM	1369	QD2	ASP	173	24.730	25.728	51.504	1.00 15.34	AAAA
MOTA	1370	С	ASP	173	25.590	23.145	53.149	1.00 16.39	AAAA
ATOM	1371	0	ASP	173 °	25.684	21.922	53.279	1.00 15.48	AAAA
	1372								
MOTA		N	GLY	174	26.583	23.912	52.705	1.00 15.58	AAAA
MOTA	1373	CA	GLY	174	27.869	23.346	52.360	1.00 13.97	AAAA
MOTA	1374	С	GLY	174	28.508	22.723	53.595	1.00 18.44	AAAA
ATOM	1375	0	GLY	174	28.970				
			177-			21.586	53.540	1.00 15.48	AAAA
ATOM	1376	N	VAL	175	28.554	23.456	54.706	1.00 16.84	AAAA
MOTA	1377	CA	ΔAT	175	29.136	22.923	55.946	1.00 16.54	AAAA
MOTA	1378	CB	VAL	175	29.201	24.031	57.037	1.00 15.88	AAAA
	1379		VAL			22.021			
MOTA				175	29.927	23.507	58.307	1.00 15.35	AAAA
ATOM	1380	CG2	VAL	175	29.923	25.258	56.476	1.00 15.62	AAAA
ATOM	1381	С	VAL	175	28.318	21.720	56.467	1.00 19.21	AAAA
ATOM	1382	ō	VAL	175	28.876	20.735	56.961		
								1.00 17.75	AAAA
ATOM	1383	N	GLN	176	26.996	21.798	56.367	1.00 17.74	AAAA
ATOM	1384	CA	GLN	176	26.164	20.685	56.832	1.00 15.66	AAAA
ATOM	1385	CB	GLN	176	24.678	20.973	56.595	1.00 16.64	AAAA
	1386	CG	GLN	176	23.789				
ATOM	100	-6	استن	110	43.109	19.788	56.952	1.00 17.00	AAAA
		•							•

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Figure 16-22

MOTA	1387	CD	GLN	176	22.325	20.106	56.884	1.00 21.52	AAAA
MOTA	1388		1 GLN	176	21.850	21.016	57.567	1.00 21.72	AAAA
	1389		GLN	176	21.581	19.348	56.064	1.00 20.30	AAAA
MOTA		C	GLN	176	26.527	19.387	56.121	1.00 16.33	AAAA
MOTA	1390			176	26.751	18.354	56.748	1.00 17.46	AAAA
MOTA	1391	0	GLN			19.443	54.799	1.00 22.24	AAAA
MOTA	1392	N	GLU	177	26.581			1.00 22.24	
MOTA	1393	CA	GLU	177	26.909	18.251	54.021		AAAA
MOTA	1394	СВ	GLU	177	26.857	18.587	52.533	1.00 15.55	AAAA
MOTA	1395	CG	GLU	177	27.131	17.388	51.623	1.00 20.24	AAAA
MOTA	1396	CD	GLU	177	26.960	17.740	50.159	1.00 27.00	AAAA
MOTA	1397	OE]	LGLU	177	27.974	17.935	49.450	1.00 30.23	AAAA
MOTA	1398	OE2	GLU	177	25.796	17.853	49.725	1.00 26.89	AAAA
MOTA	1399	С	GLU	177	28.284	17.713	54.376	1.00 20.42	AAAA
MOTA	1400	0	GLU	177	28.486	16.503	54.527	1.00 17.05	AAAA
ATOM	1401	N	ALA	178	29.233	18.626	54.527	1.00 19.67	AAAA
ATOM	1402	CA	ALA	178	30.611	18.259	54.839	1.00 18.18	AAAA
MOTA	1403	CB	ALA	178	31.464	19.519	54.918	1.00 12.76	AAAA
MOTA	1404	č	ALA	178	30.806	17.418	56.106	1.00 17.56	AAAA
	1405	ŏ	ALA	178	31.690	16.555	56.167	1.00 17.72	AAAA
MOTA	1406		PHE.		29.981	17.656	57.116	1.00 18.82	AAAA
MOTA				179	30.124	16.945	58.379	1.00 20.26	AAAA
MOTA	1407	CA	PHE	179	30.554	17.948	59.439	1.00 13.17	AAAA
MOTA	1408	CB	PHE			18.693	59.048	1.00 16.28	AAAA
ATOM	1409	CG	PHE	179	31.779			1.00 10.25	AAAA
MOTA	1410		PHE	179	31.705	20.017	58.610		AAAA
MOTA	1411		PHE	179	33.002	18.031	58.995	1.00 15.57	AAAA
MOTA	1412		PHE	179	32.845	20.673	58.114	1.00 20.03	
MOTA	1413		PHE	179	34.145	18.677	58.500	1.00 20.30	AAAA
MOTA	1414	CZ	PHE	179	34.060	20.002	58.058	1.00 19.51	AAAA
MOTA	1415	С	PHE	179	28.882	16.219	58.833	1.00 18.52	AAAA
MOTA	1416	0	PHE	179	28.773	15.828	60.000	1.00 20.21	AAAA
MOTA	1417	N	TYR	180	27.969	16.016	57.895	1.00 18.33	AAAA
ATOM	1418	CA	TYR	180	26.698	15.379	58.176	1.00 19.93	AAAA
ATOM	1419	CB	TYR	180	25.874		56.894	1.00 20.97	AAAA
MOTA	1420	CG	TYR	180	24.402	15.341	57.159	1.00 19.80	AAAA
ATOM	1421	CD1	TYR	180 ·	23.565	14.337	56.686	1.00 23.87	AAAA
ATOM	1422	CEI	TYR	180	22.203	14.391	56.898	1.00 21.32	AAAA
ATOM	1423	CD2	TYR	180	23.831	16.416	57.865	1.00 19.02	AAAA
ATOM	1424	CE2	TYR	180	22.470	16.482	58.084	1.00 26.84	AAAA
ATOM	1425	CZ	TYR	180	21.659	15.462	57.594	1.00 30.54	AAAA
ATOM	1426	OH	TYR	. 180	20.310	15.514	57.794	1.00 22.81	AAAA
ATOM	1427	c	TYR	180	26.855	13.970	58.737	1.00 22.61	AAAA
MOTA	1428	ō	TYR	180	26.064	13.526	59.579	1.00 23.44	AAAA
MOTA	1429	N	ASP	181	27.893	13.298	58.253	1.00 22.27	AAAA
ATOM	1430	CA	ASP	181	28.245		58.590	1.00 33.84	AAAA
	1431	CB	ASP	181	28.916	11.318	57.339	1.00 41.74	AAAA
MOTA	1432	CG	ASP	181	30.035	10.363	57.662	1.00 57.71	AAAA
MOTA	1433		ASP	181	30.999		58.340	1.00 61.40	AAAA
MOTA	1434		ASP	181	29.965	9.197	57.221	1.00 65.77	AAAA
ATOM		C	ASP	181	29.107	11.654	59.838	1.00 30.21	AAAA
MOTA	1435		ASP	181	29.307	10.497	60.227	1.00 27.84	AAAA
MOTA	1436	0		_		12.696			AAAA
ATOM	1437	N	THR	182	30.472	12.466	61.634	1.00 21.19	AAAA
ATOM	1438	CA	THR	182		12.977	61.358	1.00 26.55	AAAA
ATOM	1439	CB	THR	182	31.918			1.00 25.62	AAAA
MOTA	1440		THR	182	32.729	12.763	62.513		AAAA
ATOM	1441	CG2	THR	182	31.922	14.471	61.037	1.00 21.67	
ATOM	1442	С	THR	182	30.010	13.050	62.954	1.00 25.02	AAAA
ATOM	1443	0	THR	182	29.306	14.049	62.992	1.00 23.56	AAAA
MOTA	1444	N	ASP	183	30.434	12.424	64.042	1.00 19.66	AAAA
ATOM	1445	CA	ASP	183	30.086	12.894	65.371	1.00 21.52	AAAA
ATOM	1446	CB	ASP	183	29.735	11.700	66.275	1.00 28.52	AAAA
ATOM	1447	CG	ASP	183	30.920	10.783	66.523	100 32.30	AAAA
MOTA	1448		ASP	183	31.667	10.502	65.565	1.00 30.99	AAAA
ATOM	1449		ASP	183	31.095	10.326	67.675	1.00 48.65	AAAA
ATOM	1450	C	ASP	183	31.257	13.685	65.947	1.00 16.66	AAAA
ATOM	1451	ō	ASP	183	31.236	14.092	67.104	1.00 23.37	AAAA.
	1452	N	GLN	184	32.286	13.909	65.131	1.00 21.95	AAAA
MOTA	1476	44	المين		J				•

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Figure 16-23

	ATOM	1453	3 C	A GLN	184	33.43	7 14.672	65.590	1.00 17.65	AAAA
	ATOM	1454	1 C	B GLN	184	34.70				AAAA
	MOTA	1455				35.06				AAAA
	MOTA	1456				36.48		64.691	1.00 31.96	AAAA
	MOTA	1457		E1 GLN	184	36.89				AAAA
	ATOM	1458		E2 GLN		37.23				AAAA
	ATOM	1459		GLN	184	33.20				AAAA
	MOTA	1460		GLN	184	33.88				AAAA
	MOTA ATOM	1461 1462		VAL VAL	185 185	32.25				AAAA
	ATOM	1463			185	31.93 - 32.26				AAAA
	ATOM	1464		1 VAL	185	31.99				AAAA
٠	ATOM	1465		2 VAL	185	33.72				AAAA AAAA
	ATOM	1466		VAL	185	30.44			1.00 16.77	AAAA
	ATOM	1467		VAL	185	29.65				AAAA
	ATOM	1468	N	PHE		30.08			1.00 18.73	AAAA
	ATOM	1469	CA	PHE	186	28.68				AAAA
	ATOM	1470	CE	PHE	186	28.43	2 19.559	66.952		AAAA
	. ATOM	1471	CG	PHE	186	26.97		67.299	1.00 17.96	AAAA
	MOTA	1472		1 PHE	186	26.31	9 18.656	67.968	1.00 23.24	AAAA
	MOTA	1473		2 PHE	186	26.24			1.00 15.41	AAAA
	ATOM	1474		1 PHE	186	24.95				AAAA
	ATOM	1475		2 PHE	186	24.87			1.00 24.05	AAAA
	MOTA	1476	CZ		186	24.23				AAAA
	MOTA	1477 1478	C	PHE	186	28.43			1.00 17.16	AAAA
	ATOM ATOM	1479	N O	PHE VAL	186 187	29.19 27.39				AAAA
	ATOM	1480	CA		187	27.07			1.00 19.67 1.00 17.74	AAAA
	MOTA	1481	CB	VAL	187	27.01			1.00 17.74	AAAA AAAA
	ATOM	1482		1 VAL		26.57			1.00 17.31	AAAA
	ATOM	1483		2 VAL ·		28.35			1.00 16.65	AAAA
	ATOM	1484	С	VAL	187	25.73			1.00 18.46	AAAA
	ATOM	1485	0	VAL	187	24.75				AAAA
	ATOM	1486	Ņ	LEU	188	25.70	3 23.899	64.150	1.00 14.42	AAAA
	ATOM	1487	CA	LEU	188	24.48		64.567	1.00 16.68	AAAA
	ATOM	1488	CB	LEU	188	24.568		66.009	1.00 13.98	AAAA
	MOTA	1489	CG	LEU	188	23.522		66.450	1.00 13.66	· AAAA
	ATOM	1490		LEU	188	22.103			1.00 15.55	AAAA
	ATOM ATOM	1491 1492		LEU LEU	188	23.844		67.861	1.00 16.40	AAAA
	ATOM	1493	0	LEU	188 188	24.272 25.164	•		1.00 20.01	AAAA
	ATOM	1494	N	SER	189	23.10		63.506	1.00 18.86	AAAA AAAA
	ATOM	1495	CA	SER	189	22.841			1.00 14.56	AAAA
	ATOM	1496	CB	SER	189	22.896			1.00 15.55	AAAA
	ATOM	1497	OG	SER	189	22.619		60.008	1.00 14.09	AAAA
	ATOM	1498	C	SER	189	21.487		62.508	1.00 15.24	AAAA
	MOTA	1499	0	SER	189	20.509		62.578	1.00 21.46	AAAA
•	ATOM	1500	N	LEU	190	21.423		62.690	1.00 14.92	AAAA
	MOTA	1501	CA	LEU	190	20.128			1.00 15.54	AAAA
	ATOM	1502	CB	LEU	190	20.084			1.00 21.02	AAAA
	ATOM	1503 1504	CG	LEU	190	20.594		65.339	1.00 30.17	AAAA
	ATOM	1505			190	19.736			1.00 19.75	AAAA
	ATOM ATOM	1506	CDZ	LEU	190 190	20.547 20.035		65.831 61.456	1.00 19.08	AAAA
	MOTA	1507	Ö	LEU	190	21.031		60.951	1.00 14.31 1.00 15.43	AAAA
	ATOM	1508		HIS	191	18.855			1.00 15.43	AAAA AAAA
	ATOM	1509	CA	HIS	191	18.732		59.535	1.00 14.34	AAAA
	ATOM	1510	CB	HIS	191	19.506		58.539	1.00 17.34	AAAA
	ATOM	1511	CG	HIS	191	19.229		58.697	1.00 14.27	AAAA
	ATOM	1512		HIS	191	19.941		59.319	1.00 9.60	AAAA
	ATOM	1513		HIS	191	18.073	27.940	58.247	1.00 21.22	AAAA
	ATOM	1514	CE1	HIS	191	18.088		58.582	1.00 17.22	AAAA
	ATOM	1515	NE2	HIS	191	19.212		59.232	1.00 20.70	AAAA
	ATOM	1516	С	HIS	191	17.277			1.00 16.19	AAAA
	ATOM	1517	0	HIS	191	16.381		59.766	1.00 16.73	AAAA
	ATCM ·	1518	N	GLN	192	17.044	31.796	58.045	1.00 14.78	AAAA
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		^-	100	-	-		<u> </u>			
		V	تولوه	a a # 2 2 4		CET /DMI	r oel			

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MOTA	1519	CA	GLN	192	15.683	31.968	57.516	1.00 16.3	AAAA E
ATOM	1520	CB	GLN		15.669	32.871	56.283	1.00 17.0	7 AAAA
	1521	CG	GLN	192	16.174	34.270	56.498	1.00 18.1	
MOTA	1522	CD	GLN	192	16.408	34.965	55.177	1.00 14.7	
MOTA				192	15.490	35.566	54.587	1.00 20.4	
MOTA	1523		GLN			34.839	54.665	1.00 23.4	
ATOM -			GLN	192	17.630	30.584			
MOTA	1525	С	GLN	192	15.262		57.072	1.00 15.0	
MOTA	1526	0	GLN	192	16.071	29.843	56.514	1.00 19.2	
MOŢA	1527	N	SER	193	14.007	30.223	57.304	1.00 15.6	
ATOM	1528	CA	SER	193	13.561	28.907	56.877	1.00 13.8	
MOTA	1529	CB	SER	193	12.097	28.677	57.284	1.00 17.2	١
MOTA	1530	OG	SER	193	11.639	27.439	56.750	1.00 17.5	AAAA 8
ATOM	1531	С	SER	193	- 13.687	28.704	55.350	1.00 11.8	AAAA 0
ATOM	1532	0	SER	193	13.400	29.601	54.569	1.00 18.4	4 AAAA
MOTA	1533	N	PRO	194	. 14.103	27.505	54.928	1.00 14.5	9 AAAA
ATOM	1534	CD	PRO	194	14.335	26.325	55.782	1.00 19.2	2 <u>AAAA</u>
MOTA	1535	CA	PRO	194	14.268	27.143	53.513	1.00 15.3	
ATOM	1536	CB	PRO	194	14.892	25.737	53.573	1.00 18.3	
	1537	CG	PRO	194	15.359	25.587	55.007	1.00 22.3	
MOTA				194	12.880	27.104		1.00 16.4	
MOTA	1538	C	PRO	194	12.757	27.104	51.640	1.00 19.4	
ATOM	1539	0	PRO			27.151	53.681	1.00 20.5	
MOTA	1540	N	GLU	195	11.828				
ATOM	1541	CA	GLU	195	10.483	27.161	53.099	1.00 30.1	
MOTA	1542	CB	GLU	195	9.386	27.037	54.173	1.00 31.9	
ATOM	1543	CG	GLU	195	8.987	28.325	54.879	1.00 45.6	
ATOM	1544	CD	GLU	195	7.880	29.119	54.174	1.00 34.4	
ATOM	1545		GLU	195	7.635	30.259	54.612	1.00 43.9	
ATOM	1546	OE2	GLU	195	. 7.241	28.627	53.210	1.00 38.3	
ATOM	1547	С	GLU	195	10.333	28.474	52.318	1.00 26.9	
ATOM	1548	0	GLU	195	9.522	28.557	51.395	1.00 24.5	
MOTA	1549	N	TYR	196	11.116	29.501	52.669	1.00 18.1	
MOTA	1550	CA	TYR	196	11.024	30.753	51.922	1.00 15.8	
ATOM ·	1551	CB	TYR	196	10.208	31.801	52.690	1.00 20.0	1 AAAA
ATOM	1552	CG	TYR	196	10.868	32.353	53.932	1.00 19.7	7 AAAA
MOTA	1553		TYR	196	11.779	33.408	53.853	1.00 18.2	4 AAAA
ATOM	1554		TYR	196	12.407	33.898	54.988	1.00 18.5	AAAA 0
ATOM	1555		TYR	196	10.598	31.801	55.185	1.00 18.1	2 AAAA
ATOM	1556		TYR	196	11.223	32.283	56.339	1.00 21.0	AAAA e
ATOM	1557	CZ	TYR	196	12.125	33.326	56.235	1.00 20.3	AAAA e
ATOM	1558	ОН	TYR	196	12.759	33.784	57.367	1.00 16.2	
ATOM	1559	c	TYR	196	12.342	31.372	51.475	1.00 16.8	
ATOM	1560	ö	TYR	196	12.336	32.347	50.718	1.00 23.0	
	1561	N	ALA	197	13.466	30.817	51.911	1.00 17.5	
MOTA	1562	CA	ALA	197		31.400	51.512	1.00 20.2	_
ATOM		CB	ALA	197	15.315	32.261	52,659	1.00 20.7	
MOTA	1563			197	15.814	30.392	51.074	1.70 13.5	
MOTA	1564	C	ALA			29.229	51.457	1 0 19.3	
MOTA	1565	0	ALA	197	15.787	30.869	50.257	1. 70 18.0	
MOTA	1566	N	PHE	198	16.757			1.00 17.9	
MOTA	1567	CA	PHE	198	17.861	30.049	49.782		
MOTA	1568	CB	PHE	198		30.933	49.119	1.00 20.3 1.00 23.6	
ATOM	1569	CG	PHE	198	20.094	30.162	48.545	1.00 23.0	1 7777
ATOM	1570	CD1	PHE	198	20.039	29.660	47.245	1.00 29.7	1 AAAA
MOTA	1571	CD2		198	21.229	29.899	49.321	1.00 19.0	
MOTA	1572	CE1	PHE	198	21.091	28.908	46.719	1.00 30.3	
ATOM	1573	CE2	PHE	198	22.290	29.145	48.807	1.00 23.1	
ATOM	1574	CZ	PHE	198	22.218	28.646	47.493	1.00 22.7	
MOTA	1575	c	PHE	198	18.453	29.419	51.032	1.00 16.0	2 AAAA
ATOM	1576	ō	PHE	198	18.552	30.073	52.061	1.00 20.9	
ATOM	1577	N	PRO	199	18.941	28.176	50.937	1.00 19.9	2 AAA A
	1578	CD	PRO	199	19.600	27.508	52.074	1.00 17.8	
MOTA	1579	CA	PRO	199	18.990	27.318	49.744	1.00 23.5	
ATOM			PRO	199	20.108	26.344	50.095	1.00 22.7	AAAA 0
MOTA	1580	CB		199	19.813	26.087	51.534	1.00 23.1	6 AAAA
MOTA	1581	CG	PRO			26.595		1.00 30.9	7 AAAA
ATOM .		C	PRO	199	17.710		49.312	1.00 23.2	
ATOM	1583	0	PRO	199	17.733	25.855	48.322	1.00 23.2	2 AAAA
ATOM	1584	N	PHE	200	16.621	26.795	50.054	1.00 20.3	· AMMA

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ATOM	1585	CA	PHE	200		15.319	26.166	49.752	1.00	20.27	AAAA
ATOM	1586		PHE			14.840	26.533	48.346		19.77	AAAA
ATOM	1587	CG	PHE	200		14.752	27.999	48.082		18.06	AAAA
ATOM	1588	CD1	PHE	200		15.742	28.644	47.346		18.97	AAAA
MOTA	1589	CD2	PHE	200		13.654	28.736	48.519		19,06	AAAA
ATOM	1590		PHE	200		15.635	30.003	47.042		21.67	AAAA
ATOM	1591	CE2	PHE	200		13.539	30.101	48.221		22,60	AAAA
ATOM	1592	CZ	PHE	200		14.527	30.736	47.482		18.93	AAAA
ATOM	1593	С	PHE	200		15.294	24.637	49.845		18.44	AAAA
ATOM	1594	Ō	PHE	200		14.302	24.049	50.272		20.74	AAAA
ATOM	1595	N	GLU	201		16.384	24.004	49.418		20.77	AAAA
ATOM	1596	CA	GLU	201		16.522	22.542	49.399		27.34	AAAA
ATOM	1597	CB	GLU	201		17.498	22.146	48.284		28.99	AAAA
ATOM	1598	CG	GLU	201		17.024	22.458	46.881		34.82	AAAA
ATOM	1599	CD	GLU	201		18.123	22.265	45.848		32.40	AAAA
ATOM	1600	OE1	GLU	201		18.701	21.155	45.769		38.28	AAAA
ATOM	1601	OE2	GLU	201		18.405	23.230	45.111		40.08	AAAA
ATOM	1602	С	GLU	201		17.007	21.891	50.695		23.51	AAAA
ATOM	1603	0	GLU	201		16.845	20.689	50.886		23.17	AAAA
ATOM	1604	N	LYS	202		17.619	22.681	51.571	1.00	20.03	AAAA
MOTA	1605	CA	LYS	202		18.178	22.177	52.829	1.00	17.01	AAAA
MOTA	1606	CB	LYS	202		19.666	21.862	52.634	1.00	19.24	AAAA
ATOM	1607	CG	LYS	202		19.903	20.769	51.611	1.00	36.04	AAAA
ATOM	1608	CD	LYS	202		20.997	21.162	50.648	1.00	45.11	AAAA
MOTA	1609	ÇE	LYS	202		21.060	20.209	49.463	1.00	55.83	AAAA
MOTA	1610	NZ	LYS	202		22.024	20.662	48.422		28.09	AAAA
ATOM	1611	C	LYS	202		18.016	23.240	53.899		17.02	AAAA
MOTA	1612	0	LYS	202	-	17.705	24.381	53.585	1.00	20.20	AAAA
MOTA	1613	N	GLY	203		18.232	22.875	55.160		22.94	AAAA
MOTA	1614	CA	GLY	203		18.064	23.850	56.223		19.38	AAAA
MOTA	1615	C	GLY	203		16.874	23.564	57.128		20.48	AAAA
MOTA	1616	0	GLY	203		16.607	24.312	58.070		18.55	AAAA
ATOM	1617	N	PHE	204		16.150	22.484	56.852		15.42	AAAA
ATOM	1618	CA	PHE	204		14.983	22.143	57.670		20.73	AAAA
ATOM	1619	CB.	PHE	204		14.018	21.212	56.903		19.97	AAAA
MOTA	1620	CG	PHE	204		13.441	21.838	55.667		19.63	AAAA
MOTA	1621		PHE	204 204		14.137	21.801	54.459		24.96	AAAA
ATOM	1622 1623	CD2 CE1		204		12.230 13.636	22.523 22.438	55.725 53.327		18.92 20.66	AAAA
MOTA	1624	CE2		204		11.720	23.169	54.597		24.86	AAAA AAAA
ATOM ATOM	1625	CZ	PHE	204		12.422	23.127	53.400		23.66	AAAA
ATOM	1626	C	PHE	204		15.376	21.513	59.006		18.73	AAAA
ATOM	1627	ŏ	PHE	204		16.415	20.851	59.131		20.18	AAAA
ATOM	1628	N	LEU	205		14.518	21.726	59.994	1.00	19.46	AAAA
MOTA	1629	CA	LEU	205		14.727	21.244	61.356		21.09	AAAA
MOTA	1630	CB	LEU	205		13.547	21.674	62.233		23.44	AAAA
ATOM	1631	CG	LEU	205		13.506	21.222	63.693		23,23	AAAA
ATOM	1632	CD1	LEU	205		14.717	21.736	64.445	1.00	24.06	AAAA
ATOM	1633	CD2	LEU	205		12.224	21.743	64.312	1.00	30.63	AAAA
ATOM	1634	С	LEU	205		14.943	19.748	61.489	1.00	23.53	AAAA
ATOM	1635	0	LEU	205		15.659	19.315	62.381	1.00	21,28	AAAA
ATOM	1636	N	GLU	206	٠	14.356	18.959	60.591	1.00	21.59	AAAA
ATOM	1637		GLU	206		14.487	17.502	60.686	1.00	27.89	AAAA
ATOM	1638	CB	GLU	206		13.345	16.816	59.928	1.00	28.90	AAAA
MOTA	1639	CG	GLU	206		12.060	17.615	59.942	1.00	48.55	AAAA
ATOM	1640	CD	GLU	206		12.169	18.832	59.042	1.00	46.86	AAAA
ATOM	1641	OE1	GLU	206		11.360	19.769	59.178		21.58	AAAA.
ATOM	1642		GLU	206		13.076	18.833	58.181		63.58	AAAA
ATOM	1643	C	GLU	206		15.819	16.955	60.188		22.86	AAAA
MOTA	1644		GLŲ	206		16.071	15.753	60.286		21.21	AAAA
ATOM	1645	N	GLŪ	207		16.666	17.816	59.631		25.04	AAAA
ATOM	1646	CA	GLU	207		17.976	17.373	59.152		19.61	AAAA
ATOM	1647	CB (GLU	207		18.483	18.322	58.055		20.75	AAAA
ATOM	1648	CG (GLU	207		17.682	18.222	56.753		18.44	AAAA
ATOM	1649	CD (GLU	207		17.687	19.514	55.983		24.75	AAAA
MOTA	1650	OE1	GLU	207		18.738	20.182	55.948	1.00	22.17	AAAA

ATOM	165	1 01	E2 GLU	207	16.646	19.854	55.396	1.00 15.50	AAAA
ATOM	1652		. GTA		18.921	17.379	60.350		AAAA
ATOM	1653		GLU		19.506	18.416			
ATOM	1654		ILE		19.081	16.218	60.988		AAAA
MOTA	1655				19.930	16.138	62.168	1.00 21.60	AAAA
MOTA	1656			208	19.113	15.652	63.403	1.00 22.37	AAAA
MOTA	1657		2 ILE	208	19.968			1.00 28.84	AAAA
						15.693	64.653	1.00 43.26	AAAA
MOTA	1658			208	17.905	16.561	63.625	1.00 21.61	AAAA
ATOM	1659		1 ILE	.208	17.029	16.160	64.786	1.00 41.40	AAAA
ATOM	1660		ILE	208	21.156	15.260	61.981	1.00 24.74	AAAA
ATOM	1661		ILE	208	21.785	14.850	62.943	1.00 22.68	AAAA
MOTA	1662		GLY	209	21.512	14.969	60.738	1.00 22.80	AAAA
MOTA	1663			209	22.690	14.153	60.535	1.00 20.43	AAAA
MOTA	1664	C	GLY	209	22.342	12.769	60.037	1.00 25.56	AAAA
MOTA	1665	0	GLY	209	21.165	12.447	59.850		AAAA
MOTA	1666	N	GLU	210	23.373	11.944	59.888	1.00 26.07	AAAA
MOTA	1667	CA	GLU-	210	23.235	10.601	59.348	1.00 25.78	AAAA
MOTA	1668	CB	GLU	210	23.404	10.731	57.835	1.00 28.27	AAAA
ATOM	1669	CG	GLU	210	23.041	9.569	56.965	1.00 56.41	AAAA
ATOM	1670	CD	GLU	210	23.170	9.952	55.495	1.00 65.37	AAAA
MOTA	1671	OE	1 .GLU	210	24.290	10.327	55.075	1.00 62.00	AAAA
MOTA	1672	OE	2 GLU	210	22.153	9.894	54.768	1.00 73.24	AAAA
ATOM	1673		GLU	210	24.329	9.709	59.936	1.00 31.85	AAAA
ATOM	1674	0	GLU	210	25.447	10.170	60.217	1.00 28.85	AAAA
ATOM	1675	N	GLY	211	24.012	8.431	60.121	1.00 27.84	AAAA
ATOM	1676	CA	GLY	211	24.991	7.502	60.657	1.00 26.25	AAAA
ATOM	1677	C	GLY	211	25.545	7.942	61.995	1.00 27.79	AAAA
ATOM	1678	ō	GLY	211	24.788	8.324	62.874	1.00 28.66	AAAA
ATOM	1679	N	LYS	212	26.865	7.880	62.150	1.00 34.62	AAAA
ATOM	1680	CA	LYS.	212	27.512	8.287	63.393	1.00 34.39	AAAA
ATOM	1681	СВ	LYS	212	29.029	8.132	63.273	1.00 40.40	AAAA
ATOM	1682	CG	LYS	212	29.505	6.712	62.996	1.00 53.97	
ATOM	1683	CD	LYS	212	29.139	5.770	64.131		AAAA
MOTA	1684	CE	LYS	212	29.612	4.347	63.863	1.00 61.93 1.00 62.74	AAAA
ATOM	1685	NZ	LYS	212	31.091	4.258	63.711		AAAA
MOTA	1686	C	LYS	212	27.181	9.741		1.00 70.11	AAAA
ATOM	1687	ò	LYS	212	27.109	10.126	63.725	1.00 36.04	AAAA
MOTA	1688	N	GLY	213	26.959	10.543	64.897	1.00 28.34	AAAA
	1689	CA	GLY	213	26.648	11.948	62.688	1.00 31.47	AAAA
MOTA	1690	C	GLY	213	25.189	12.291	62.898	1.00 31.68	AAAA
MOTA	1691	Ö	GLY	213	24.840		63.142	1.00 28.78	AAAA
ATOM	1692	N	LYS	214		13.460	63.259	1.00 22.56	AAAA
MOTA	1693	CA			24.317	11.292	63.222	1.00 28.54	AAAA
MOTA	1694		LYS	214	22.905	11.585	63.463	1.00 31.11	AAAA
ATOM		CB	LYS	214	22.080	10.295	63.325	1.00 31.03	AAAA
ATOM	1695	CG	LYS	214	20.583	10.461	63.224	1.00 38.15	AAAA
ATOM	1696	CD	LYS	214	19.968	9.115	62.844	1.00 40.49	AAAA
ATOM	1697	CE	LYS	214	18.490	9.220	62.537	1.00 48.02	AAAA
MOTA	1698	NZ	LYS	214	17.927	7.924	62.064	1.00 44.99	AAAA
MOTA	1699	C	LYS	214	22.834	12.160	64.875	1.00 26.90	AAAA
MOTA	1700	0	LYS	214	23.260	11.524	65.831	1.00 33.33	AAAA
MOTA	1701	N	GLY	215	22.310	13.376	64.997	1.00 24.38	AAAA
MOTA	1702	CA	GLY	215	22.230	14.034	66.290	1.00 26.03	AAAA
ATOM	1703	С	GLY	215	23.298	15.115	66.447	1.00 27.03	AAAA
MOTA	1704	0	GLY	215	23.352	15.820	67.458	1.00 23.34	AAAA
MOTA	1705	N	TYR	216	24.152	15.260	65.439	1.00 22.79	AAAA
MOTA	1.706	CA	TYR	216	25.217	16.257	65.512	1.00 25.51	AAAA
ATOM	1707	CB	TYR	216	26.592	15.576	65.406	1.00 20.54	AAAA
MOTA	1708	CG	TYR	216	26.900	14.671	66.581	1.00 26.47	AAAA
MOTA	1709		TYR	216	26.221	13.464	66.757	1.00 28.96	AAAA
ATOM	1710	CE1	TYR	216	26.455	12.660	67.872	1.00 33.08	AAAA
MOTA	1711	CD2	TYR	216	27.832	15.052	67.552	1.00 24.21	AAAA
MOTA	1712	CE2	TYR	216	28.074	14.254	68.675	1.00 34.45	AAAA
ATOM	1713	CZ	TYR	216	27.378	13.063	68.827	1.00 40.53	AAAA
ATOM	1714	OH	TYR	216	27.580	12.291	69.947	1.00 45.67	AAAA
ATOM	1715	С	TYR	216	25.104	17.391	64.493	1.00 22.57	AAAA
ATOM	1716	0	TYR	216		18.014	64.126	1.00 19.70	AAAA
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MOTA	1717	N	ASN	217	23.889	17.635	64.027	1.00 2	2.88	AAAA
ATOM	1718	CA	ASN	217	23.621	18.729	63.109	1.00 2	2.60	AAAA
			ASN	217	23.453	18.240	61.671	1.00 1		AAAA
ATOM	1719	CB								
MOTA	1720	CG	ASN	217	23.233	19.387	60.695	1.00 1		AAAA
MOTA	1721	OD1	ASN	217	22.098	19.704	60.307	1.00 2		AAAA
MOTA	1722	ND2	ASN	217	24.320	20:032	60.309	1.00 1	.2.18	AAAA
ATOM	1723.	C	ASN	217	22.311	19.296	63.630	1.00 1	7.65	AAAA
				217	21.381	18.550	63.894	1.00 1		AAAA
MOTA	1724	0	ASN							
MOTA	1725	N	LEU	218	22.236	20.610	63.793	1.00 2		AAAA
MOTA	1726	CA	LEU	218	21.014	21.197	64.320	1.00 2	11.20	AAAA
MOTA	1727	CB	LEU	218 -	21.186	21.547	65.808	1.00 1	.7.73	AAAA
	1728	CG	LEU	218	19.906	21.702	66.647	1.00 3	2.30	AAAA
MOTA					20.228	22.427	67.944	1.00 2		AAAA
MOTA	1729		LEU	218						
MOTA	1730	CD2	LEU	218	18.862	22.464	65.903	1.00 4		AAAA
MOTA	1731	С	LEU	218	20.700	22.459	63.554	1.00 1		AAAA
ATOM	1732	0	LEU	218	21.467	23.425	63.615-	1.00 1	.6.70	AAAA
	1733	N	ASN	219	19.590	22.441	62.824	1.00 1	5.43	AAAA
MOTA				219	19.143	23.609	62.072	1.00 1		AAAA
MOTA	1734	CA	ASN							
MOTA	1735	CB	ASN	219	18.634	23.232	60.665	1.00 1		AAAA
ATOM	1736	CG	ASN	219	19.732	22.738	59.750	1.00 2		AAAA
ATOM	1737	OD1	ASN	219	20.861	23.232	59.802	1.00 1	.7.90	'AAAA
ATOM	1738		ASN	219	19.398	21.789	58.868	1.00 1	6.62	AAAA
	1739		ASN	219	17.990	24.256	62.821	1.00 2	1.98	AAAA
MOTA		C				23.569	63.262	1.00 1		AAAA
MOTA	1740	0	ASN	219	17.075					
MOTA	1741	N	ILE	220	18.025	25.580	62.952	1.00 1		AAAA
MOTA	1742	CA	ILE	220	16.951	26.298	63.640	1.00 1		AAAA
MOTA	1743	CB	ILE	220	17.522	27.115	64.823	1.00 1	L5.70	AAAA
ATOM	1744		ILE	220	16.411	27.912	65.479	1.00 1	5.18	AAAA
	1745		ILE	220	18.246	26.193	65.823	1.00 1	9.11	AAAA
MOTA				220	17.350	25.259	66.632	1.00 2		AAAA
ATOM -	1746		ILE					1.00 1		AAAA
MOTA	1747	С	ILE	220	16.363	27.246	62.573			
ATOM	1748	0	ILE	220	16.810	28.386	62.419	1.00 1		AAAA
MOTA	1749	N	PRO	221	15.341	26.790	61.826	1.00 1		AAAA
MOTA	1750	CD	PRO	221 ·	14.612	25.518	61.906	1.00 1	L8.83	AAAA
ATOM	1751	CA	PRO	221.	14.739	27.628	60.785	1.00 1	L9.83	AAAA
	1752	CB	PRO	221	13.930	26.615	59.948	1.00 2	20.76	AAAA
MOTA			PRO	221	14.409	25.241	60.462	1.00 2		AAAA
ATOM	1753	CG				28.664	61.444	1.00 2		AAAA
MOTA	1754	С	PRO	221	13.849					
ATOM	1755	0	PRO	221	13.061	28.318	62.314	1.00 2		AAAA
ATOM	1756	N	LEU	222	13.977	29.926	61.028	1.00 1		AAAA
ATOM	1757	ÇA	LEU	222	13.209	31.018	61.612	1.00 2		AAAA
MOTA	1758	CB	LEU	222	14.163	31.972	62.319	1.00 1	L6.46	AAAA
ATOM	1759	CG	LEU	222	14.868	31.232	63.466	1.00 1	L8.65	AAAA
		CD1		222	16.026	32.072	64.014	1.00.2	21.32	AAAA
MOTA	1760				13.857	30.925	64.555	1.00 1		AAAA
ATOM	1761	CD2		222				1.00 1		AAAA
MOTA	1762	C	LEU	222	12.350	31.763	60.590			
ATOM	1763	0	LEU	222	12.687	31.830	59.412	1.00 1		AAAA
MOTA	1764	N	PRO	223	11.220	32.329	61.042	1.00 1	•	AAAA
ATOM	1765	CD	PRO	223	10,723	32.249	62.431	1.00 1	17.38	AAAA
	1766	CA	PRO	223	10.264	33.065	60.203	1.00 1	19.59	AAAA
MOTA		CB	PRO	223	9.006	33.083	61.074	1.00 2		AAAA
MOTA	1767						62.441			AAAA.
MOTA	1768	CG	PRO	223	9.608	33.304				
ATOM	1769	С	PRO	223	10.606	34.458	59.723	1.00 2		AAAA
ATOM	1770	0	PRO	223	11.525	35.101	60.214	1.00 1		AAAA
MOTA	1771	N	LYS	224	9.830	34.912	58.7 45	1.00 1	16.41	AAAA
ATOM	1772	CA	LYS	224	9.975	36.254	58.200	1.00 1	L6.11 '	AAAA
	1773	CB	LYS	224	9.002	36.446	57.039	1.00 2	20.34	AAAA
MOTA					9.163	35.441	55.900	1.00 1		AAAA
MOTA	1774	CG	LYS	224				1.00 2		AAAA
ATOM	1775	CD	LYS	224	8.109	35.687	54.807			
MOTA	1776	CE	LYS	224	8.209	34.624	53.704	1.00 2		AAAA
ATOM	1777	NZ	LYS	224	7.207	34.843	52.618	1.00		AAAA
MOTA	1778		LYS	224	9.638	37.289	59.284	1.00 1	L5.77	AAAA
	1779	ō	LYS	224	8.819	37.032	60.186	1.00 2	21.07	AAAA
ATOM			GLY	225	10.239	38.469	59.171	1.00 2		AAAA
	1780	N		225	9.974	39.527	60.129	1.00 2		AAAA
ATOM	1781	CA	GLY		J.J/4			1.00 2	20.63	AAAA
ATOM	1782	C	GLY	225	10.556	.39.286	61.507	1.00 4		-

						20 012	CD 460	1.00 20.66	AAAA
ATOM	1783	0	GLY	225	10.128	39.912	62.468	1.00 20.37	AAAA
MOTA	1784	N	LEU	226	11.540	38.395	61.606		
ATOM	1785	CA	LEU	226	12.154	38.063	62.893	1.00 18.71	AAAA
ATOM	1786	CB	LEU	226	13.354	37.145	62.670	1.00 13.63	AAAA
ATOM	1787	CG	LEU	226	13.836	36.443	63.939	1.00 18.44	AAAA
ATOM -	1788	CD1		226	12.834	35.329	64.243	1,00 18.09	AAAA
	1789	CD2		226	15.232	35.844	63.741	1.00 17.96	AAAA
MOTA	1790	C	LEU	226	12.649	39.309	63.642	1.00 19.84	AAAA
MOTA		ō	LEU	226	13.320	40.151	63.052	1.00 18.13	AAAA
MOTA	1791			227	12.336	39.421	64.932	1.00 23.30	AAAA
MOTA	1792	N	ASN			40.571	65.692	1.00 20.88	AAAA
MOTA	1793	CA	ASN	227	11.682	41.261	66.485	1.00 21.73	AAAA
MOTA	1794	CB	ASN	.227			67.546		AAAA
MOTA	1795	CG	ASN	227	11.061	40.368	68.341	1.00 23.80	AAAA
MOTA	1796	OD1	ASN	227	11.762	39.736		1.00 21.08	AAAA
MOTA	1797	ND2	ASN	227	9.729	40.340	67.581		
ATOM	1798	С	ASN	227	13.950	40.152	66.612	1.00 25.24	AAAA-
MOTA	1799	0	ASN	227	14.282	38.965	66.702	1.00 18.54	AAAA
ATOM	1800	N	ASP	228	14.547	41.124	67.296	1.00 19.41	AAAA
ATOM	1801	CA	ASP	228	15.682	40.844	68.169	1.00 22.15	AAAA
ATOM	1802	CB	ASP	228	16.208	42.141	68.802	1.00 16.82	AAAA
ATOM	1803	CG	ASP	228	16.852	43.060	67.796	1.00 30.68	. AAAA
	1804		ASP	228	17.182	42.576	66.690	1.00 23.87	AAAA
ATOM	1805		ASP	228	17.053	44.256	68.123	1.00 25.02	AAAA
MOTA		C	ASP	228	15.440	39.835	69.265	1.00 18.83	AAAA
MOTA	1806		ASP	228	16.298	39.002	69.536°	1.00 16.28	AAAA
MOTA	1807	0		229	14.291	39.930	69.928	1.00 20.73	AAAA
MOTA	1808	N	ASN		13.975	39.015	71.007	1.00 21.75	AAAA
MOTA	1809	CA	ASN	229	12.706	39.483	71.712	1.00 19.46	AAAA
ATOM	1810	CB	ASN	229		40.738		1.00 27.14	AAAA
MOTA	1811	CG	ASN	229	12.943	40.691	73.556	1.00 33.03	AAAA
ATOM	1812		ASN	229	13.588		72.019	1.00 21.35	AAAA
MOTA	1813		ASN	229	12.464	41.874 37.596	70.503	1.00 18.47	AAAA
MOTA	1814	С	ASN	229	13.833		70.303	1.00 22.47	AAAA
MOTA	1815	0	ASN	229	14.284	36.644.		1.00 17.79	AAAA
ATOM	1816	N	GLU	230	13.252	37.454	69.319	1.00 17.73	AAAA
ATOM	1817	ÇA	GLU	230	13.081	36.125	68.748		AAAA
ATOM	1818	CB	GLU	、230	12.152	36.193	67.536	1.00 20.54	AAAA
ATOM	1819	CG	GLU	230	10.765	36.714	67.890	1.00 28.98	
ATOM	1820	CD	GLU	230	9.870	36.816	66.677	1.00 24.35	AAAA
ATOM	1821	OE1	GLU	230	10.360	37.296	65.638	1.00 22.00	AAAA
MOTA	1822	OE2	GLU	230	8.683	36.443	66.772	1.00 24.99	AAAA
ATOM	1823	С	GLU	230	14.422	35.507	68.361	1.00 16.89	AAAA
ATOM	1824	Ó	GLU	230	14.663	34.326	68.603	1.00 19.45	AAAA
ATOM	1825	N	PHE	231	15.305	36.305	67.772	1.00 15.68	AAAA
	1826	CA	PHE	231	16.616	35.788	67.389	1.00 15.78	AAAA
MOTA	1827	CB	PHE	231	17.420	36.863	66.649	1.00 13.22	AAAA
MOTA	1828	CG	PHE	231	18.719	36.361	66.069	1 00.20.63	AAAA
MOTA			PHE	231	18.723	35.445	65.016	1 00 18.42	AAAA
MOTA	1829		PHE	231	19.936	36.804	66.568	1 70 21.10	AAAA
MOTA	1830			231	19.918	34.983	64.471	1.00 17.67	AAAA
MOTA	1831		PHE	231	21.144	36.346	66.029	1.00 28.29	AAAA
MOTA	1832		PHE		21.130	35.431	64.976	1.00 27.85	` AAAA
MOTA	1833	CZ		231	17.385	35.332	68.636	1.00 18.54	AAAA
MOTA	1834	С	PHE	231	17.363	34.201	68.702	1.00 18.86	AAAA
MOTA	1835	0	PHE	231	17.869	36.204	69.636	1.00 19.07	AAAA
MOTA	1836	N	LEU	232	17.495		70.848	1.00 17.39	AAAA
ATOM	1837	CA	LEU	232	18.239	35.850		1.00 24.53	AAAA
MOTA	1838	CB	LEU	232	18.415	37.078	71.737	1.00 16.64	AAAA
ATOM	1839	CG	LEU	232	19.214	38.202	71.061	1.00 16.04	AAAA
MOTA	1840	CD1	LEU	232	19.134	39.449	71.934	1.00 20.70	
MOTA	1841		LEU	232	20.659	37.806	70.810	1.00 18.77	AAAA
MOTA	1842	c	LEU	232	17.607	34.707	71.628	1.00 19.82	AAAA
	1843	õ	LEU	232	18.309	33.904	72.217	1.00 21.80	AAAA
ATOM	1844	N	PHE	233	16.281	34.640	71.648	1.00 17.18	AAAA
MOTA		CA	PHE	233	15.587	33.537	72.309	1.00 23.34	AAAA
ATOM	1845	CB	PHE	233	14.074	33.663	72.095	1.00 19.17	AAAA
MOTA	1846		PHE	233	13.289	32.447	72.523	1.00 21.40	AAAA
ATOM	1847	CG			12.863	32.302	73.841	1.00 29.62	AAAA
MOTA	1848	CDI	PHE	233		J 2 . J 4 D		•	•

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ATOM	1849	CD2	PHE	233	12.942	31.473	71.596	1.00 19.92	AAAA
	1850	CE1	PHE	233	12.088	31.206	74.229	1.00 29.35	
ATOM									AAAA
MOTA	1851	CE2	PHE	233	12.168	30.363	71.966	1.00 25.37	AAAA
MOTA	1852	CZ	PHE	233	11.737	30.231	73.283	1.00 30.28	AAAA
ATOM	1853	С	PHE	233	16.041	32.234	71.660	1.00 23.12	AAAA
ATOM	1854	0	PHE	233	16.433	31.273	72.332	1.00 18.35	AAAA
MOTA	1855	N	ALA	234	15.961	32.208	70.332	1.00 17.26	AAAA
ATOM	1856	CA	ALA	234	16.332	31.026	69.562	1.00 17.67	AAAA
ATOM	1857	CB	ALA	234	16.085	31.297	68.046	1.00 19.08	AAAA
MOTA	1858	С	ALA	234	17.786	30.641	69.800	1.00 16.31	AAAA
ATOM	1859	0	ALA	234	18.127	29.461	69.926	1.00 16.75	AAAA
	1860	N	LEU	235	18.646	31.643	69.846	1.00 16.73	AAAA
ATOM									
MOTA	1861	CA	LEU	235	20.074	31.411	70.051	1.00 19.14	AAAA ·
MOTA	1862	CB	LEU	235	20.823	32.742	69.956	1.00 21.72	AAAA
					22.226	32.790	69.345	1.00 36.73	
MOTA	1863	CG	LEU	235					AAAA
ATOM	1864	CD1	LEU	235	23.026	33.844	70.105	1.00 20.69	AAAA
MOTA	1865	CDS	LEU	235	22.917	31.426	69.393	1.00 22.96	AAAA
	_								
MOTA	1866	С	LEU	235	20.354	30.776	71.421	1.00 18.71	AAAA
MOTA	1867	0	LEU	235	21.028	29.747	71.522	1.00 18.59	AAAA
	1868	N	GLU	236	19.831	31.390	72.479	1.00 25.43	AAAA
MOTA									
ATOM	1869	CA	GLU	236	20.046	30.883	73.839	1.00 19.75	AAAA
MOTA	1870	CB	GLU	236	19.335	31.777	74.860	1.00 23.18	AAAA
		CG			19.725	33.229	74.777	1.00 38.53	
MOTA	1871		GLU	236					AAAA
MOTA	1872	CD	GLU	236	18:857	34.119	75.648	1.00 42.42	AAAA
MOTA	1873	OE1	GLU	236	17.617	34.171	75.428	1.00 45.43	AAAA
			GLU	236	19.425	34.768	76.548	1.00 48.76	
ATOM	1874	OE2					•		AAAA
ATOM	1875	С	GLU	236	19.541	29.452	74.011	1.00 25.85	AAAA
ATOM	1876	0	GLU	236	20,222	28.603	74.597	1.00 21.36	AAAA
	1877	Ŋ	LYS	237	18.343	29.193	73.501	1.00 23.16	AAAA
MOTA									
ATOM	1878	CA	LYS	237	17.752	27.871	73.610	1.00 17.06	AAAA
ATOM	1879	CB	LYS	237	16.282	27.943	73.193	1.00 26.98	AAAA
ATOM	1880	CG	LYS	237	15.483	26.711	73.519	1.00 52.00	AAAA
						27.110		1.00 56.40	
MOTA	1881	CD	LYS	237	14.078		73.932		AAAA
ATOM	1882	CE	LYS	237	14.131	27.979	75.183	1.00 52.03	AAAA
ATOM	1883	NZ	LYS	237	12.782	28.421	75.614	1.00 55.53	AAAA
	1884	C	LYS	237	18.502	26.827	72.785	1.00 18.46	AAAA
MOTA									
ATOM	1885	0	LYS	237	18.691	25.692	73.231	1.00 21.20	AAAA
ATOM	1886	N	SER	238	18.932	27.187	71.578	1.00 21.28	AAAA
ATOM	1887	CA	SER	238	19.649	26.208	70.776	1.00 16.47	AAAA
					19.745		69.307	1.00 19.75	AAAA
ATOM	1888	CB	SER	238		26.666			
ATOM	1889	OG	SER	238	20.475	27.858	69.160	1.00 22.52	AAAA
MOTA	1890	С	SER	238	21.039	25.923	71.361	1.00 18.79	AAAA
		Õ	SER	238	21.521	24.788	71.312	1.00 20.60	AAAA
MOTA	1891								
ATOM	1892	N	LEU	239	21.690	26.937	71.925	1.00 22.95	AAAA
ATOM	1893	CA	LEU	239	23.004	26.701 [.]	72.513	1.00 2098	AAAA
	1894	CB.	LEU	239	23.652	28.008	72.986	1.00 18.39	AAAA
ATOM									
MOTA	1895	CG	LEU	239	23.985	29.072	71.933	1.00 20.02	AAAA
ATOM	1896	CD1	LEU	239	24.538	30.311	72.636	1.00 27.02	AAAA
ATOM	1897	CD2	T.ETT	239	25.010	28.556	70.933	1.00 20.31	AAAA
MOTA	1898	С	LEU	239	22.882	25.735	73.680	1.00 25.16	AAAA
ATOM	189 9	0	LEU	239	23.780	24.929	73.920	1.00 20.70	AAAA
ATOM	1900	N	GLU	240	21.768	25.800	74.398	1.00 24.93	AAAA
						24.912	75.536		
ATOM	1901	CA	GLU	240	21.570			1.00 25.72	AAAA
ATOM	1902	CB	GLU	240	20.331	25.356	76.337	1.00 29.10	AAAA
ATOM	1903	CG	GLU	240	20.042	24.531	77.581	1.00 49.56	AAAA
							78.515	1.00 63.15	AAAA
MOTA	1904		GLU	240	19.053	25.212			
ATOM	1905	OE1	GLU	240	17.935	25.550	78.067	1.00 69.26	AAAA
ATOM	1906	OE2	GLU	240	19.400	25.410	79.703	1.00 66.68	AAAA
			GLU	240	21.440	23.469	75.046	1.00 23.44	AAAA
MOTA	1907								
ATOM	1908	0	GLU	240	21.951	22.535	75.674	1.00 23.10	AAAA
ATOM	1909	N	ILE	241	20.771	23.294	73.913	1.00 19.52	AAAA
	1910		ILE	241	20.598	21.978	73.321	1.00 24.06	AAAA
ATOM									
MOTA	1911		ILE	241	19.705	22.039	72.052	1.00 23.80	AAAA
ATOM	1912	CG2	ILE	241	19.718	20.678	71.323	1.00 24.94	AAAA
	1913	CG1		241	18.281	22.433	72.439	1.00 28.60	AAAA
ATOM									
MOTA	1914	CD1	TTE.	241	17.336	22.600	71.257	1.00 27.04	AAAA
		•		•				-	•

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					•					
MOTA	1915	С	ILE	241	21.	957 21.	404 72.9	41 1.00	25.48	AAAA
ATOM	1916	ō	ILE		22.		234 73.1		19.43	AAAA
	1917	N	VAL		22.		235 72.3		20.41	AAAA
MOTA					24.		782 71.9		23.17	
ATOM	1918	CA	VAL							AAAA
MOTA	1919	CB	VAL		24.		856 71.1		26.48	AAAA
MOTA	1920		L VAL		26.		394 70.8		18.67	AAAA
MOTA	1921	CG2	: VAL	242	24.	093 23.	135 69.8		26.97	AAAA
ATOM	1922	С	VAL	242	24.	962 21.	456 73.1	54 1.00	24.81	AAAA
ATOM	1923	0	VAL	242	25.	566 20.	384 73.2	35 1.00	22.49	AAAA
ATOM	1924	N	LYS		24.5	989 22.	387 74.1	02 1.00	26.06	AAAA
ATOM	1925	CA	LYS		25.				32.57	AAAA
	1926	CB	LYS		25.9		379 76.2		28.53	AAAA
ATOM			LYS		26.				43.21	AAAA
MOTA	1927	CG					191 78.6		53.10	AAAA
MOTA	1928	CD	LYS		26.0					
ATOM	1929	CE	LYS	243	26.4				50.30	AAAA
MOTA	1930	NZ	LYS	243	26.0				59.15	AAAA
MOTA	1931	C	LYS	243	25.4				30.38	AAAA
MOTA	1932	0	LYS	243	26.3	321 20.	255 76.5 °	78 1.00	35.44	AAAA
ATOM	1933	N	GLU	244	24.3	61 20.	542 76.0	76 1.00	28.12	AAAA
ATOM	1934	CA	GLU	244	23.7	798 19.	320 76.7	98 1.00	37.54	AAAA
ATOM	1935	CB	GLU	244	22.2	88 19.	260 77.0	48 1.00	35.34	AAAA
ATOM	1936	CG	GLU	244	21.7				55.88	
	1937	CD	GLU	244	20.2				57.89	AAAA
ATOM			GLU	244	19.6				60.60	AAAA
ATOM	1938				19.7				57.73	AAAA
MOTA	1939		GLU	244					38.17	
ATOM	1940	C	GLU	244	24.2					AAAA
MOTA	1941	0	GLU	244	24.2				38.46	AAAA
MOTA	1942	N	VAL	245	. 24.5				30.29	AAAA
MOTA	1943	CA	VAL	245	24.9				29.17	AAAA
ATOM	1944	CB	VAL	245	23.9	84 16.			46.68	AAAA
MOTA	1945	CG1	VAL	245	24.4	62 15.	641 71.9	42 1.00	53.09	AAAA
ATOM	1946	CG2	VAL	245	22.5	81 16.	488 73.3	27 1.00	54.19	AAAA
ATOM	1947	С	VAL	245	26.3	64 16.	982 73.5	08 1.00	34.90	AAAA
MOTA	1948	ō	VAL	245	26.9				34.73	AAAA
MOTA	1949	N	PHE	246	26.9				29.22	AAAA
	1950	CA	PHE	246	28.3				29.17	AAAA
MOTA			PHE	246	28.1				30.42	AAAA
MOTA	1951	CB							25.62	AAAA
MOTA	1952	CG	PHE	246	29.3				28.89	
MOTA	1953		PHE	246	29.6					AAAA
MOTA	1954		PHE	246	30.1				25.17	AAAA
MOTA	1955		PHE	246	30.7				23.43	AAAA
MOTA	1956	CE2	PHE	246	31.2				22.40	AAAA
MOTA	1957	CZ	PHE	246	31.5	49 18.	236 68.8	35 1.00	19.88	AAAA
MOTA	1958	С	PHE	246	29.2	33 19.	176 73.7	12 1.00	23.38	AAAA
MOTA	1959	0	PHE	246	28.8	67 20.	312 74.0	02 1.00	29.15	AAAA
ATOM	1960	N	GLU	247	30.4	10 18.	682 74.0	94 1.00	29.73	AAAA
ATOM	1961	CA	GLU	247	31.3				28.10	AAAA
	1962	СВ	GLU	247	31.9				35.75	AAAA
MOTA				247	30.9				60.78	AAAA
ATOM	1963	CG	GLU						70.07	
MOTA	1964	CD	GLU	247	29.7				79.95	AAAA
MOTA	1965	OE1		247	28.9					
MOTA	1966		GLU	247	29.4			35 1.00	76.80	AAAA
MOTA	1967	С	GLU	247	32.5				28.90	AAAA
ATOM	1968	0	GLU	247	33.4	90 18.	946 73.7		23.67	AAAA
ATOM	1969	N	PRO	248	32.5	31 20.	391 73.18	B1 1.00	25.02	AAAA
ATOM	1970	CD	PRO	248	31.5	74 22.0	003 73.3	10 1.00	27.23	AAAA
ATOM	1971	CA	PRO	248	33.5			09 1.00	28.06	AAAA
	1972	CB	PRO	248	33.0				28.11	AAAA
MOTA		CG	PRO	248	31.5				34.57	AAAA
MOTA	1973								23.87	AAAA
MOTA	1974	C	PRO	248	34.9				24.05	AAAA
ATOM	1975	0	PRO	248	35.1					
MOTA	1976	N	GLU	249	35.9				24.34	AAAA
ATOM	1977	CA	GLU	249	37.3			1.00	25.98	AAAA
	1978	CB	GLU	249	38.2			79 1.00	22.07	AAAA
ATOM	1979	CG	GLU	249	38.0	46 18.1			33.40	AAAA
ATOM	1980	CD	GLU	249	39.0	05 17.3	767 71.4	15 1.00	29.15	AAAA
	-						-			

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Figure 16-31

ATOM.	1981	OE.	1 GLU	249	39.071	17.770	70.199	1.00 27.62	222
ATOM	1982		2 GLU	249	39.694			1.00 27.82	AAAA AAAA
ATOM	1983		GLU	249	37.692		71.786	1.00 26.04	AAAA
ATOM	1984	0	GLU	249	38.582		72.262	1.00 26.39	AAAA
ATOM	1985	N	VAL	250	36.953		70.744	1.00 23.83	AAAA
ATOM	1986	CA	VAL	250	37.151	24.197	70.086	1.00 19.67	AAAA
MOTA	1987	CB	VAL	250	38.438		69.210	1.00 20.88	AAAA
MOTA	1988	CG:	1 VAL	250	38.348	23.117	68.128	1.00 18.18	AAAA
ATOM	1989	CG	2 VAL	-250	38.647	25.530	68.591	1.00 16.71	AAAA
ATOM	1990	С	VAL	250	35.946	24.483	69.207	1.00 20.78	AAAA
ATOM .	1991	0	VAL	250	. 35.299	23.556	68.746	1.00 19.60	AAAA
MOTA	1992	N	TYR	251	35.633	25.757	69.000	1.00 18.75	AAAA
ATOM	1993	CA	TYR	251	34.497	26.109	68.153	1.00 22.44	AAAA
MOTA	1994	CB	TYR	251	33.261	26.437	69.022	1.00 16.57	AAAA
MOTA	1995 1996	CG CD1	TYR	251	33.207	27.856	69.575	1.00 22.36	AAAA
MOTA MOTA	1997	CE		251 251	32.654 32.612	28.896 30.185	68.823	1.00 18.12	AAAA
ATOM	1998	CD2		251	33.715	28.160	69.308 70.842	1.00 20.40	AAAA
ATOM	1999	CE2		251	33.676	29.475	71.349	1.00 20.04 1.00 16.60	AAAA
ATOM	2000	CZ	TYR	251	33.128	30.473	70.573	1.00 14.68	AAAA AAAA
ATOM	2001	OH	TYR	251	33.100	31.780	71.011	1.00 21.79	AAAA
ATOM	2002	C	TYR	251	34.811	27.294	67.236	1.00 20.28	AAAA
ATOM	2003	O	TYR	251	35.695	28.107	67.525	1.00 19.91	AAAA
ATOM	2004	N	LEU	252	34.097	27.360	66.109	1.00 17.90	AAAA
ATOM	2005	CA	LEU	252	34.216	28.466	65.161	1.00 18.58	AAAA
MOTA	2006	CB	LEU	252	34.679	28.001	63.767	1.00 17.55	AAAA
MOTA	2007	CG	LEU	252	36.028	27.290	63.718	1.00 23.36	AAAA
MOTA	2008		LEU	252	35.819	25.820	64.017	1.00 27.78	AAAA
ATOM	2009		LEU	252	36.631	27.440	62.331	1.00 27.29	AAAA
ATOM	2010	Ç	LEU	252	32.816	29.049	65.052	1.00 15.49	AAAA
ATOM	2011	0	LEU ·		31.819	28.320	65.120	1.00 18.82	AAAA
MOTA MOTA	2012 2013	N CA	LEU LEU	253 253	32.756 31.498	30.360 31.105	64.891	1.00 16.80	AAAA
ATOM	2014	CB	LEU	253	31.379	31.987	64.817 66.073	1.00 17.50 1.00 15.49	AAAA AAAA
MOTA	2015	ĊĠ	LEU	253	30.326	33.085	66.165	1.00 17.75	AAAA
ATOM	2016		LEU	253	28.946	32.438	66.172	1.00 20.85	AAAA
ATOM	2017		LEU	253	30.536	33.897	67.464	1.00 19.05	AAAA
ATOM	2018	С	LEU	253	31.516	31.985	63.580	1.00 20.22	AAAA
MOTA	2019	0	LEU	253	32.474	32.727	63.371	1.00 18.14	AAAA
MOTA	2020	N	GLN	254	30.466	31.913	62.765	1.00 16.50	AAAA
ATOM	2021	CA	GLN	254	30.411	32.730	61.556	1.00 16.48	AAAA
MOTA	2022	CB	GLN	254	30.085	31.863	60.312	1.00 25.58	AAAA
MOTA	2023	CG	GLN	254	28.647	31.798	59.871	1.00 36.40	AAAA
ATOM	2024 2025	CD OE1	GLN	254	28.337	32.728	58.701	1.00 33.18	AAAA
ATOM ATOM	2025	NE2	GLN GLN	254 254	28.744 27.613	32.487 33.799	57.546 58.992	1.00 21.05 1.00 22.85	AAAA
ATOM	2027	C	GLN	254	29.384	33.799	61.832	1.00 22.85	AAAA AAAA
ATOM	2028=	ŏ	GLN	254	28.282	33.577	62.364	1.00 10.12	AAAA
ATOM	2029	N	LEU	255	29.768	35.032	61.468	1.00 14.42	AAAA
ATOM	2030	CA	LEU	255	28.988	36.215	61.763	1.00 17.99	AAAA
ATOM	2031	CB	LEU	255	29.834	37.070	62.719	1.00 20.68	AAAA
MOTA	2032	CG	LEU	255	30.240	36.283	63.964	1.00 22.90	AAAA
ATOM	2033	CD1	LEU	255	31.446	36.906	64.635	1.00 29.36	AAAA
ATOM	2034	CD2	LEU	255	29.042	36.214 ·		1.00 14.80	AAAA
ATOM	2035	Ç	LEU	255	28.541	37.060	60.594	1.00 19.32	AAAA
ATOM	2036		LEU	255	28.838	38.260	60.561	1.00 21.23	AAAA
MOTA	2037	N	GLY	256	27.827	36.467	59.639	1.00 17.21	AAAA
ATOM	2038	CA	GLY	256	27.347	37.259	58.516	1.00 15.64	AAAA
ATOM	2039	C	GLY	256	26.413	38.348	59.028	1.00 17.31	AAAA
MOTA	2040	O N	GLY	256 257	25.717	38.150	60.027 58.348	1.00 15.62	AAAA
ATOM	2041 2042	N CA	THR THR	257 257	26.389 25.536	39.494 40.598	58.776	1.00 19.72 1.00 19.88	AAAA
ATOM ATOM	2042	CB	THR	257	26.242	40.538	58.589	1.00 19.88	AAAA AAAA
ATOM	2044	OG1		257	26.538	42.187	57.206	1.00 14.02	AAAA
ATOM	2045		THR	257	27.543	42.009	59.392	1.00 17.58	AAAA
ATOM	2046		THR	257	24.199	40.634	58.053	1.00 20.58	AAAA
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Figure 16-32

ATOM	2047	0	THE	257	23.403	41.545	58.266	1.00 14.59	AAAA
ATOM	2048		ASP		23.927	39.639	57.213	1.00 16.56	
	2049				22.651	39.646		•	
MOTA								1.00 16.39	
MOTA	2050				22.604	38.611	55.388	1.00 18.38	
MOTA	2051	CG			23.037	37.229	55.811	1.00 25.85	
ATOM -			1 ASP		23.222	36.995	57.022	1.00 22.32	
MOTA	2053	OD:	2 ASP	258	23.187	36.370	54.909	1.00 18.12	AAAA
MOTA	2054	С	ASP	25B	21.396	39.563	57.397	1.00 21.25	AAAA
MOTA	2055	0	ASP	258	20.300	39.781	56.897	1.00 22.52	
ATOM	2056	N	PRO		21.510	39.172	58.680	1.00 18.17	
ATOM	2057	CD	PRO		22.614	38.528	59.422	1.00 25.88	
ATOM	2058	CA	PRO		20.281	39.139	59.482	1.00 21.24	
ATOM	2059	CB			20.710			- 1.00 21.18	
-			PRO			38.363			
MOTA	2060	CG	PRO	259	22.174	38.707	60.846	1.00 36.11	
ATOM	2061	С	PRO		19.705	40.534	59.820	1.00 20.88	
ATOM	2062	0	PRO		18.572	40.647	60.280	1.00 19.25	
MOTA	2063	N	LEU	260	20.473	41.591	59.571	1.00 18.75	AAAA
ATOM	2064	CA	LEU	260	20.023	42.949	59.875	1.00 22.16	AAAA
ATOM	2065	CB	LEU	260	21.202	43.935	59.778	1.00 20.35	AAAA
ATOM	2066	CG	LEU	260	22.403	43.640	60.682	1.00 21.82	AAAA
MOTA	2067	CD1	LEU	260	23.604	44.486	60.253	1.00 18.57	
ATOM	2068		LEU	260	22.032	43.873	62.123	1.00 19.18	
MOTA	2069	C	LEU	260	18.876	43.469	59.014	1.00 24.16	
MOTA	2070	ò	LEU	260	18.742	43.144	57.826	1.00 21.69	
							59.634		
ATOM	2071	N	LEU	2.61	18.049	44.300		1.00 19.54	
MOTA	2072	CA	LEU	261	16.903	44.913	58.965	1.00 17.34	
ATOM	2073	CB	LEU	261	16.285	45.967	59.892	1.00 19.96	
ATOM	2074	CG	LEU	261	15.204	46.879	59.300	1.00 29.99	
MOTA	2075		LEU	261	14.080	46.040	58.732	1.00 33.66	
MOTA	2076		LEU	261	14.682	47.819	60.376	1.00 44.71	
ATOM	2077	С	LEU	261	17.262	45.550	57.620	1.00 18.11	
MOTA	2078	0	LEU	261	16.539	45.386	56.634	1.00 19.02	AAAA
ATOM	2079	N	GLU	262	18.391	46.249	57.566	1.00 22.68	AAAA
ATOM	2080	CA	GLU	262 .	18.802	46.921	56.338	1.00 18.46	AAAA
ATOM	2081	CB	GLU	262	19.875	47.965	56.641	1.00 22.01	
MOTA	2082	CG	GLU	262	19.365	49.136	57.443	1.00 22.94	
ATOM	2083	CD	GLU	262	19.434	48.902	58.927	1.00 23.11	
ATOM	2084		GLU	262	19.668	47.748	59.357	1.00 24.58	
ATOM	2085		GLU	262	19.238	49.883	59.667	1.00 27.06	
	2086	C	GLU	262	19.281	46.034	55.197	1.00 25.65	
ATOM	_							1.00 25.49	
ATOM	2087	0	GLU	262	19.446	46.510	54.070		
MOTA	2088	И	ASP	263	19.501	44.750	55.467	1.00 22.45	
MOTA	2089	CA	ASP	263	19.959	43.851	54.418	1.00 15.93	
ATOM	2090	CB	ASP	263	20.981	42.859	54.988	1.00 18.99	
ATOM	2091	CG	ASP	263	21.706	42.081	53.907	1.00 22.21	
MOTA	2092		ASP	263	22.876	41.730	54.139	1.00 23.19	
MOTA	2093	OD2	ASP	263	21.112	41.809	52.838	1.00 25.02	
MOTA	2094	C	ASP	263	18.733	43.165	53.837	1.00 22.32	AAAA
MOTA	2095	0	ASP	263	18.012	42.419	54.519	1.00 18.50	AAAA
ATOM	2096	N	TYR	264	18.500	43.447	52.564	1.00 25.21	AAAA
MOTA	2097	CA	TYR	264	17.339	42.936	51.865	1.00 29.92	AAAA
ATOM	2098	CB	TYR	264	17.077	43.776	50.596	1.00 38.48	AAAA
ATOM	2099	CG	TYR	264	17.910	43.431	49.379	1.00 54.09	AAAA
	2100		TYR	264	17.677	42.249	48.660	1.00 69.38	AAAA
ATOM			TYR	264			47.526	1.00 68.71	AAAA
MOTA	2101				18.420	41.930			
ATOM	2102	CD2	TYR	264	18.915	44.286	48.928	1.00 66.09	
MOTA	2103		TYR	264	19.670	43.975	47.788	1.00 74.50	AAAA
MOTA	2104	CZ	TYR	264	19.415	42.794	47.094	1.00 72.57	AAAA
ATOM	2105	ОН	TYR	264	20.154	42.472	45.975	1.00 71.96	AAAA
MOTA	2106	С	TYR	264	17.445	41.461	51.532	1.00 29.55	AAAA
ATOM	2107	0	TYR	264	16.448	40.839	51.190	1.00 30.11	AAAA
ATOM	2108	N	LEU	265	18.639	40.891	51.629	1.00 24.45	AAAA
ATOM	2109	CA	LEU	265	18.753	39.476	51.337	1.00 25.36	AAAA
ATOM	2110	CB	LEU	265	20.186	39.089	50.969	1.00 29.81	AAAA
		CG	LEU	265	20.509	39.510	49.531	1.00 34.43	
ATOM	2111				21.847				
MOTA	2112	CD1	i-eu	265	41.04/	38.930	49.100	1.00 44.38	AAAA

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ATOM	2113	CD2	LEU	265	19.422	38.990	48.603	1.00 46.72	AAAA
ATOM	2114	C	LEU	265	18.209	38.585	52.447	1.00 22.33	AAAA
MOTA	2115	0	LEU	265	18.279	37.364	52.348	1.00 23.48	AAAA
ATOM	2116	N	SER	266	17.677	39.194	53.508	1.00 17.50	AAAA
MOTA	2117	CA	SER	266	17.055	38.398	54.569	1.00 19.69	AAAA
MOTA	2118	CB	SER	266	17.912	38.314	55.845	1.00 20.73	AAAA
MOTA	2119	OG	SER	266	17.696	39.442	56.684	1.00 22.81	AAAA
MOTA	2120	С	SER	266	15.739	39.048	54.950	1.00 19.75	AAAA
MOTA	2121	0	SER	266	15.572	40.265	54.840	1.00 23.66	AAAA
ATOM	2122	N	LYS	267	14.799	38.229	55.402	1.00 18.40 1.00 20.64	AAAA
ATOM	2123	CA	LYS	267	13.527 12.397	38.759 37.787	55.851 55.513	1.00 20.84	AAAA AAAA
MOTA	2124 2125	CB CG	LYS LYS	267 267	12.397	37.787	54.025	1.00 25.60	AAAA '
ATOM ATOM	2125	CD	LYS	267	12.095	38.823	53.259	1.00 23.00	AAAA
ATOM.	2127	CE	LYS	267	11.985	38.540	51.772	1.00 38.49	AAAA
ATOM	2128	NZ	LYS	267	11.954	39.793	50.991	1.00 33.11	AAAA
ATOM	2129	C	LYS	267	13.601	38.987	57.365	1.00 20.63	AAAA
ATOM	2130	0	LYS	267	12.584	39.192	58.017	1.00 25.38	AAAA
MOTA	2131	N	PHE	268	14.814	38.937	57.915	1.00 18.98	AAAA
ATOM	2132	CA	PHE	268	15.034	39.182	59.345	1.00 18.50	AAAA
MOTA	2133	CB	PHE	268	16.328	38.510	59.833	1.00 20.91	AAAA
MOTA	2134	CG	PHE	268	16.252	37.006	59.967	1.00 16.96	AAAA
MOTA	2135		PHE	268	17.374	36.290	60.415	1.00 16.61	AAAA
ATOM	2136		PHE	268	15.081	36.303	59.682	1.00 18.13	AAAA
MOTA	2137		PHE	268	17.331	34.904	60.581	1.00 14.81 1.00 17.45	AAAA AAAA
MOTA	2138 2139	CEZ	PHE PHE	268 268	15.027 16.144	34.900 34.208	59.849 60.296	1.00 17.43	AAAA
ATOM ATOM	2140	C	PHE	268	15.179	40.699	59.510	1.00 18.33	AAAA
ATOM	2141	ō	PHE	268	15.733	41.371	58.644	1.00 18.28	AAAA
ATOM	2142	N	ASN	269	14.679	41.236	60.613	1.00 21.04	AAAA
ATOM	2143	CA	ASN	269	14.763	42.675	60.859	1.00 22.89	AAAA
ATOM	2144	CB	ASN	269	13.365	43.298	60.940	1.00 20.55	AAAA
MOTA	2145	CG	ASN	269	12.551	43.071.		1.00 26.13	AAAA
MOTA	2146		ASN	269	13.060	43.192	58.571	1.00 29.17	AAAA
MOTA	2147		ASN	269	11.268	42.767	59.860	1.00 28.26	AAAA
MOTA	2148	C	ASN	269	15.493	42.967	62.159	1.00 19.00	AAAA AAAA
ATOM	2149	0	ASN	269	14.984 16.695	43.683 42.435	63.019 62.298	1.00 21.85 1.00 17.71	AAAA
ATOM	2150 2151	N CA	LEU	270 270	17.441	42.642	63.521	1.00 17.71	AAAA
ATOM ATOM	2152	CB	LEU	270	18.441	41.507	63.712	1.00 18.95	AAAA
MOTA	2153	CG	LEU	270	17.945	40.058	63.631	1.00 20.54	AAAA
ATOM	2154	CD1		270	19.070	39.174	64.152	1.00 14.19	AAAA
ATOM	2155	CD2		270	16.679	39.853	64.465	1.00 19.05	AAAA
ATOM	2156	С	LEU	270	18.203	43.971	63.583	1.00 22.83	AAAA
ATOM	2157	0	LEU	270	18.409	44.643	62.560	1.00 18.25	AAAA
MOTA	2158	N	SER	271	18.621	44.318	64.799	1.00 20.95	AAAA
ATOM	2159	CA	SER	271	19.414	45.518	65.081	1.00 18.28	AAAA
ATOM	2160	CB	SER	271	18.985 19.347	46.150	66.409 67.512	1.00 18.73 1.00 22.28	AAAA AAAA
MOTA	2161	og	SER SER	271 271	20.875	45.327 45.073	65.224	1.00 19.98	AAAA
ATOM ATOM	2162 2163	С 0	SER	271	21.122	43.899	65.537	1.00 18.82	AAAA
ATOM	2164	N	ASN	272	21.828	45.994	65.020	1.00 16.17	AAAA
ATOM	2165	CA	ASN	272	23.270	45.695	65.145	1.00 20.70	AAAA
ATOM	2166	CB	ASN	272	24.176	46.903	64.884	1.00 37.49	AAAA
ATOM	2167	CG	ASN	272	24.161	47.378	63.483	1.00 54.53	AAAA
ATOM	2168	OD1	ASN	272	24.702	48.454	63.199	1.00 45.61	AAAA
MOTA	2169	ND2	ASN	272	23.576	46.594	62.579	1.00 60.55	AAAA
MOTA	2170	С	ASN	272	23.586	45.343	66.580	1.00 18.03	AAAA
ATOM	2171	0	ASN	272	24.545	44.625	66.854	1.00 18.58	AAAA
ATOM	2172	N	VAL	273	22.831	45.938	67.500	1.00 19.57	AAAA
ATOM	2173	CA	VAL	273	23.053	45.698	-68.919	1.00 22.12 1.00 26.91	AAAA AAAA
ATOM	2174	CB	VAL	273	22.345 22.440	46.765 46.421	69.765 71.233	1.00 26.51	AAAA
ATOM	2175	CG1 CG2		273 273	23.034	48.421	69.531	1.00 34.73	AAAA
MOTA	2176 2177	CG2	VAL	273	22.636	44.295	69.341	1.00 22.06	AAAA
MOTA. MOTA	2178	0	VAL	273		.43.708	70.217	1.00 16.89	AAAA .
A TOTAL		_							

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MOTA	2179	N	ALA	274	21.601	43.747	68.713	1.00 21.79	AAAA
ATOM	2180	CA	ALA	274	21.207	42.383	69.035	1.00 21.31	AAAA
ATOM	2181	CB	ALA		19.806	42.092	68.475	1.00 18.95	AAAA
ATOM	2182	C	ALA		22.259	41.451	68.400	1.00 17.83	AAAA
	2183	ō	ALA		22.569	40.389	68.947	1.00 20.38	AAAA
MOTA						41.859	67.245	1.00 16.01	
MOTA	2184	N	PHE		22.798				AAAA
MOTA	2185	CA	PHE		23.828	41.089	66.536	1.00 16.46	AAAA
MOTA	2186	CB	PHE		24.220	41.835	65.253	1.00 24.56	AAAA
MOTA	2187	CG	PHE		25.363	41.222	64.492	1.00 23.01	AAAA
MOTA	2188	CD1	PHE	275	25.209	40.035	63.788	1.00 23.88	AAAA
MOTA	2189	CD2	PHE	275	26.590	41.877	64.443	1.00 22.40	AAAA
ATOM	2190	CE1	PHE	275	26.266	39.510	63.038	1.00 28.74	AAAA
MOTA	2191	CE2	PHE	275	27.654	41.365	63.700	1.00 35.03	AAAA
MOTA	2192	CZ	PHE	275	27.489	40.181	62.996	1.00 24.63	AAAA
MOTA	2193	С	PHE	275 ·	25.030	40.964	67.469	1.00 25.06	AAAA
ATOM	2194	ō	PHE		25.619	39.888	67.632	1.00 19.71	AAAA
ATOM	2195	N	LEU		25.366	42.080	68.101	1.00 17.49	AAAA
ATOM	2196	CA	LEU		26.482	42.139	69.030	1.00 24.23	AAAA
	2197	CB	LEU	276	26.736	43.606	69.416	1.00 20.44	AAAA
ATOM	2198	CG	LEU	276	28.001	43.967	70.211	1.00 39.65	AAAA
ATOM			LEU	276	27.948	45.447	70.589	1.00 29.65	AAAA
ATOM	2199						71.460	1.00 32.41	
MOTA	2200		LEU	276	28.102	43.143			AAAA
MOTA	2201	С	LEU	276	26.180	41.278	70.262	1.00 18.88	AAAA
MOTA	2202	0	LEU	276	27.045	40.529	70.727	1.00 17.99	AAAA
MOTA	2203	N	LYS	277	24.968	41.374	70.805	1.00 19.67	AAAA
ATOM	2204	CA	LYS	277	24.644	40.552	71.964	1.00 21.33	AAAA
MOTA	2205	CB	LYS	277	23.265	40.888	72.532	1.00 23.84	AAAA
MOTA	2206	CG	LYS	277	23.247	42.126	73.366	1.00 40.87	AAAA
ATOM	2207	CD	LYS	277	22.069	42.086	74.325	1.00 54.73	AAAA
MOTA	2208	CE	LYS	277	22.172	40.884	75.254	1.00 58.85	AAAA
MOTA	2209	NZ	LYS	277	21.051	40.844	76.228	1.00 55.34	AAAA
MOTA	2210	С	LYS	277	24.695	39.068	71.660	1.00 22.12	AAAA
ATOM	2211	0	LYS	277	25.074	38.264	72.513	1.00 22.19	AAAA
ATOM	2212	N	ALA	278	24.311	38.700	70.441	1.00 20.23	AAAA
ATOM	.2213	CA	ALA	278	24.325	37.291	70.039	1.00 17.06	AAAA
ATOM	2214	CB	ALA	278	23.798	37.154	68.589	1.00 19.27	AAAA
ATOM	2215	С	ALA	278	25.760	36.767	70.127	1.00 16:94	AAAA
MOTA	2216	0	ALA	278	26.035	35.676	70.648	1.00 14.93	AAAA
ATOM	2217	N	PHE	279	26.679	37.564	69.606	1.00 18.88	AAAA
ATOM	2218	CA	PHE	279	28.099	37.231	69.626	1.00 21.01	AAAA
ATOM	2219	CB	PHE	279	28,880	38.392	68.998	1.00 16.79	AAAA
MOTA	2220	CG	PHE	279	30.370	38.264	69.120	1.00 20.23	AAAA
ATOM	2221		PHE	279	31.062	37.272	68.423	1.00 21.61	AAAA
ATOM	2222		PHE	279	31.088	39.159	69.905	1.00 23.24	AAAA
	2223		PHE	279	32.461	37.185	68.509	1.00 30.98	AAAA
ATOM	2224		PHE	279	32.480	39.081	69.995	1.00 24.82	AAAA
ATOM		CZ	PHE	279	33.169	38.095	69.295	1.00 30.27	AAAA
ATOM	2225			279	28.576	36.995	71.067	1.00 25.48	AAAA
ATOM	2226	C	PHE				71.362	1.00 16.30	AAAA
MOTA	2227	0	PHE	279	29.275	36.016		1.00 22.30	AAAA
MOTA	2228	N	ASN	280	28.194	37.898	71.962		
MOTA	2229	CA	ASN	280	28.599	37.777	73.352	1.00 24.49	AAAA
ATOM	2230	CB	ASN	280	28.391	39.109	74.080	1.00 27.17	AAAA
MOTA	2231	CG	ASN	280	29.344	40.183	73.578	1.00 20.88	AAAA
ATOM	2232	OD1	ASN	280	30.503	39.897	73.273	1.00 22.95	AAAA
MOTA	2233	ND2	ASN	280	28.875	41.421	73.522	1.00 27.85	AAAA
ATOM	2234	C	ASN	280	27.928	36.636	74.095	1.00 23.01	AAAA
ATOM	2235	0	ASN	280	28.510	36.062	75.016	1.00 21.91	AAAA
ATOM	2236	N	ILE	281	26.711	36.300	73.689	1.00 18.74	AAAA
ATOM	2237	CA	ILE	281	26.005	35.179	74.294	1.00 18.37	AAAA
ATOM	2238	СВ	ILE	281	24.566	35.067	73.758	1.00 19.31	AAAA
ATOM	2239	CG2		281	23.977	33.725	74.135	1.00 28.87	AAAA
ATOM	2240	CG1		281	23.710	36.206	74.308	1.00 23.51	AAAA
	2241	CD1		281	22.279	36.193	73.776	1.00 26.47	AAAA
MOTA	2241		ILE	281	26.743	33.876	73.965	1.00 18.54	AAAA
MOTA		C	ILE	281	26.830	32.973	74.801	1.00 19.69	AAAA
MOTA	2243	0	VAL	282	27.258		72.744	1.00 17.72	AAAA
ATOM	2244	N	AWD.	202	41.430	33.765	16.144	A	1-1-1-1

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	0045	~~		000	07 076	20 552	TO 250	1 00 14 00	
MOTA	2245	CA	.VAL	282	27.976	32.553	72.352	1.00 14.89	AAAA
ATOM	2246	CB	VAL	282	28.359	32.565	70.852	1.00 18.50	AAAA
ATOM	2247		L VAL		29.342	31.440	70.567	1.00 20.73	AAAA
MOTA	2248	CGZ	VAL		27.105	32.363	69.994	1.00 17.49	AAAA
MOTA	2249	С	VAL	282	29.241	32.433	73.198	1.00 21.79	AAAA
ATOM	2250	0	VAL		29.568	31.360	73.715	1.00 25.80	AAAA
MOTA	2251	N	ARG	283	29.935	33.549	73.361	1.00 19.14	AAAA
ATOM	2252	· CA	ARG	283	31.161	33.548	74.150	1.00 23.51	AAAA
						34.898			
MOTA	2253	CB	ARG		31.851		74.023	1.00 20.64	AAAA
ATOM	2254	CG	ARG	283	32.338	35.200	72.607	1.00 19.65	AAAA
ATOM	2255	CD	ARG	283	32.754	36.645	72.474	1.00 25.70	AAAA
					33.970	36.944			
ATOM	2256	NE	ARG				73.215	1.00 36.05	AAAA
MOTA	2257	CZ	ARG	283	34.277	38.147	73.681	1.00 34.61	AAAA
ATOM	2258	NH1	. ARG	283	33.448	39.169	73.488	1.00 35.23	AAAA
	2259		ARG		35.419	38.332	74.326		
MOTA								1.00 29.30	AAAA
MOTA	2260	С	ARG	283	30.911	33.219	75.622	1.00 25.44	AAAA
ATOM	2261	0	ARG	283	31.754	32.600	76.272	1.00 23.12	AAAA
	2262	Ň	GLU		29.765	33.632	76.151		
MOTA								1.00 26.79	AAAA
MOTA	2263	CA	GLU	284	29.462	33.338	77.553	1.00 31.77	AAAA
ATOM	2264	CB	GLU	284	28.243	34.115	78.033	1.00 30.96	AAAA
ATOM	2265	CG	GLU	284	28.399	35.605	77.957	1.00 50.56	AAAA
MOTA	2266	CD	GLU	284	27.137	36.320	78.365	1.00 63.75	AAAA
MOTA	2267	OE1	GLU	284	26.085	36.067	77.738	1.00 68.93	AAAA
			GLU	284		37.133			
ATOM	2268				27.198		79.309	1.00 72.01	AAAA
MOTA	2269	С	GLU	284	29.181	31.862	77.733	1.00 31.57	AAAA
ATOM	2270	0	GLU	284	29.410	31.310	78.803	1.00 33.08	AAAA
		N	VAL		28.673	31.221	76.686		
MOTA	2271			285				1.00 23.37	AAAA
ATOM '	2272	CA	VAL	285	· 28.354	29.807	76.774	1.00 23.25	AAAA
ATOM	2273	CB	VAL	285	27.221	29.407	75.789	1.00 24.77	AAAA
	2274	CG1	VAL	285	26.952	27.913	75.881	1.00 26.98	AAAA
ATOM									
MOTA	2275	CG2	VAL	285	25.940	30.181	76.107	1.00 24.98	AAAA
MOTA	2276	С	VAL	285	29.567	28.942	76.479	1.00 31.41	AAAA
ATOM	2277		VAL	285	29.833	27.983	77.195	1.00 25.34	AAAA
ATOM	2278	N	PHE	286	30.316		. 75.431	1.00 27.27	AAAA
ATOM	2279	CA	PHE	286	31.463	28.457	75.086	1.00 22.47	AAAA
ATOM	2280	CB	PHE	286	31.289	27.904	73.667	1.00 22.26	AAAA
ATOM	2281	CG	PHE	286	30.168	26.918	73.536	1.00 25.71	AAAA
MOTA	2282	CD1	PHE	286	28.971	27.274	72.917	1.00 22.88	AAAA
ATOM	2283	CD2	PHE	286	30.294	25.631	74.069	1.00 24.49	AAAA
ATOM .	2284		PHE	286	27.919	26.365	72.829	1.00 19.85	AAAA
ATOM	2285	CE2	PHE	286	29.246	24.714	73.987	1.00 27.48	AAAA
ATOM	2286	CZ	PHE	286	28.056	25.081	73.367	1.00 24.59	AAAA
							75.225		
MOTA	2287	С	PHE	286	32.854	29.059		1.00 21.53	AAAA
MOTA	2288	0	PHE	286	33.849	28.417	74.873	1.00 27.12	AAAA
ATOM	2289	N	GLY	287	32.937	30.272	75.754	1.00 23.76	AAAA
	2290	CA	GLY	287	34.237	30.896	75 901	1.00 24.17	AAAA
ATOM									
ATOM	2291	С	GLY	287	34.705	31.419	74.562	1.00 27.05	AAAA
ATOM	2292	0	GLY	287	33.888	31.670	•73.667	1.00 18.06	AAAA
ATOM	2293	N	GLU	288	36.017	31.576	74.414	1.00 23.21	AAAA
MOTA	2294	CA	GLU	288	36.583	32.085	73.170	1.00 24.87	AAAA
ATOM	2295	CB	GLU	288	37.968	32.682	73.410	1.00 29.25	AAAA
	2296	CG	GLU	288	37.984	33.933	74.291	1.00 42.63	
ATOM									AAAA
ATOM	2297	CD	GLU	288	37.114	35.052	73.745	1.00 43.77	AAAA
ATOM	2298	OE1	GLU	288	37.235	35.380	72.544	1.00 36.82	AAAA
	2299	OE2		288	36.317	35.617	74.521	1.00 51.56	AAAA
ATOM									
MOTA	2300	С	GLU	288	36.693	31.028	72.072	1.00 20.85	AAAA
ATOM	2301	0	GLU	288	36.995	29.856	72.332	1.00 18.10	AAAA
•	2302	N	GLY	289	36.447	31.468	70.843	1.00 26.12	AAAA
ATOM					_				
ATOM	2303	CA	GLY	289	36.517	30.588	69.692	1.00 20.71	AAAA
ATOM	2304	С	GLY	289	37.126	31.318	68.510	1.00 18.56	AAAA
	2305	ō	GLY	289	37.669	32.404	68.679	1.00 16.59	AAAA
MOTA								_	
MOTA	2306	N	VAL	290	37.032	30.724	67.322	1.00 19.86	AAAA
MOTA	2307	CA	VAL	290	37.572	31.312	66.103	1.00 19.70	AAAA
	2308		VAL	290	38.150	30.192	65.184	1.00 19.04	AAAA
ATOM									
MOTĄ	2309	CG1		290	38.667	30.769	63.853	1.00 15.54	AAAA
ATOM	2310	CG2	VAL	290	39.296	29.483 ·	65.920	1.00 20.40	AAAA
		-		•			-		

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•		_							
MOTA	2311	С	VAL	290	36.408	32.040	65.427	1.00 20.90	AAAA
ATOM	2312	0	VAL	290	35.351	31.439	65.193	1.00 19.33	AAAA
MOTA	2313	N	TYR	291	36.598	33.325	65.125	1.00 15.37	AAAA
ATOM	2314	CA	TYR	291	35.543	34.140	64.524	1.00 16.79	AAAA
						_		_	
ATOM	2315	CB	TYR		35.412		65.317	1.00 16.42	AAAA
MOTA	2316	CG	TYR	291	35.375	35.181	66.808	1.00 18.60	AAAA
			L TYR		36.366				
MOTA	2317		-				67.659	1.00 21,77	AAAA
ATOM	2318	CE:	l TYR	291	36.368	35.385	69.030	1.00 22.55	AAAA
ATOM	2319	CD2	2 TYR	. 291	34.388	34.374	67.361	1.00 17.34	
									AAAA
MOTA	2320	CE2	TYR	291	34.381	34.066	68.718	1.00 20.24	AAAA
MOTA	2321	CZ	TYR	291	- 35.367	34.568	69.545	1.00 25.85	AAAA
ATOM	2322	ОН	TYR	291	35.338	34.246	70.885	1.00 25.57	. AAAA
ATOM	2323	С	TYR	291	35.720	34.446	63.031	1.00 14.97	AAAA
ATOM	2324	0	TYR	291	36.773	34.921	62.586	1.00 15.21	
									AAAA
ATOM	2325	N	LEU	292	34.660	34.189	62.273	1.00 14.06	AAAA
MOTA	2326	CA	LEU	292	34.674	34.392	60.824	1.00 15.03	AAAA
MOTA	2327	CB	LEU	292	34.461	33.046	60.108	1.00 13.66	AAAA
ATOM	2328	CG	LEU	292	35.342	31.856	60.496	1.00 19.04	AAAA
ATOM	2329		LEU	292	34.909	30.615	59.665	1.00 15.17	
									AAAA
MOTA	2330	CD2	LEU	292	36.792	32.190	60.252	1.00 19.18	AAAA
ATOM	2331	С	LEU	292	33.564	35.327	60.396	1.00 16.62	AAAA
MOTA	2332	0	LEU	292	32.575	35.488	61.107	1.00 14.76	AAAA
ATOM	2333	N	GLY	293	33.724	35.932	59.216	1.00 18.62	AAAA
ATOM	2334	CA	GLY	293	32.696	36.816	58.699	1.00 17.10	AAAA
				293					
MOTA	2335	C	GLY		.31.611	35.954	58.068	1.00 23.44	AAAA
MOTA	2336	С	GLY	293	31.407	34.798	58.459	1.00 23.60	AAAA
MOTA	2337	N	GLY	294	30.915	36.501	57.085	1.00 24.96	AAAA
ATOM	2338	CA	GLY	294	29.871	35.738	56.434	1.00 27.07	
				-					AAAA
MOTA	2339	C	GLY	294	29.132	36.632	55.474	1.00 28.41	AAAA
ATOM	2340	0	GLY	294	29.605	37.722	55.167	1.00 25.66	AAAA
MOTA	2341	N	GLY-	295	27.972	36.168	55.011	1.00 20.33	AAAA
MOTA	2342	CA	GLY	295	27.164	36.936	54.085	1.00 20.14	AAAA
ATOM	2343	C	GLY	295	26.742	38.244	54.730	1.00 25.34	AAAA
ATOM	2344	.0	GLY	295	26.550	38.317	55.942	1.00 28.89	AAAA
		•							
MOTA	2345	N	GLY	296	26.614	39.274	53.909	1.00 28.52	AAAA
ATOM	2346	CA	GLY	296	26.230	40.598	54.367	1.00 23.21	AAAA
ATOM	2347	С	GLY	296	26.314	41.342	53.059	1.00 26.34	AAAA
	2348			296	27.359	41.324			
MOTA		0	GLY				52.414	1.00 26.05	AAAA
MOTA	2349	N	TYR	297	25.235	42.008	52.662	1.00 22.61	AAAA
MOTA	2350	CA	TYR	297	25.228	42.644	51.360	1.00 22.58	AAAA
	2351	СВ	TYR	297	24.265	41.861	50.457	1.00 23.68	
ATOM									AAAA
MOTA	2352	CG	TYR	29 7	24.502	40.352	50.521	1.00 25.14	AAAA
MOTA	2353	CD1	TYR	297	23.981	39.571	51.568	1.00 28.31	AAAA
	2354	CE1	TYR	297	24.269	38.196	51.662		AAAA
ATOM								1.00 24.18	
ATOM	2355	CD2	TYR	297	25.307	39.725	49.577	1.00 29.74	AAAA
MOTA	2: 36	CE2	TYR	297	25.598	38.362	49.664	1.00 27.09	AAAA
ATOM	257	CZ	TYR	297	25.085	37.606	50.696	1.00 28.68	AAAA
ATOM	8ذذ2	OH	TYR	297	25.407	36.261	50.739	1.00 28.17	AAAA
ATOM	2359	С	TYR	297	24.916	44.138	51.320	1.00 24.98	AAAA
	2360	0	TYR	297	24.841	44.714	50.237	1.00 26.51	AAAA
ATOM	_								
MOTA	2361	N.	HIS	298	24.740	44.752	52.491	1.00 23.80	AAAA
MOTA	2362	CA	HIS	298	24.480	46.188	52.591	1.00 23.44	AAAA
MOTA	2363	CB ·		298	23.325	46.494	53.536	1.00 23.37	AAAA
MOTA	2364	CG	HIS	298	22.956	47.945	53.551	1.00 32.94	AAAA
ATOM	2365	CD2	HIS	298	23.491	48.983	54.232	1.00 24.31	AAAA
	2366	ND1		298	22.011	48.487	52.707	1.00 38.29	AAAA
MOTA									
ATOM	2367	CE1	HIS	298	21.978	49.797	52.868	1.00 26.60	AAAA
ATOM	2368	NE2	HIS	298	22.867	50.125	53.788	1.00 36.57	AAAA
	2369	C	HIS	298	25.757	46.775	53.184	1.00 23.11	AAAA
MOTA									
MOTA	2370	0	HIS	298	26.135	46.439	54.306	1.00 22.64	AAAA
ATOM	2371	N	PRO	299	26.430	47.673	52.445	1.00 21.07	AAAA
			PRO	299	26.078	48.207	51.117	1.00 27.42	AAAA
ATOM	2372	CD							
ATOM	2373	CA	PRO	299	27.676	48.286	52.910	1.00 26.62	AAAA
ATOM ·	.2374	CB	PRO	299	28.041	49.228	51.755	1.00 28.95	AAAA
		CG	PRO	299	26.678	49.600	51.196	1.00 35.16	AAAA
ATOM	2375								
ATOM	2376	С	PRO	299	27.644	48.991	54.262	1.00 25.75	AAAA

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MOTA	2377	0	PRO	299	28.565	48.845	55.068	1.00 24.36	AAAA
	2378	N	TYR						
MOTA			_		26.602		54.504	1.00 24.48	
ATOM	2379	CA	TYR	300	26.495	50.478	55.766	1.00 22.94	AAAA
ATOM	2380	CB	TYR	300	25.317		55.734	1.00 25.24	
MOTA	2381	CG	TYR	300	25.411	52.599	54.762	1.00 30.44	AAAA
MOTA	2382	CD1	L TYR	300	26.366	52.634	53.746	1.00 26.01	. AAAA
MOTA	2383		L TYR		26.389	53.676	52.819	1.00 29.66	
ATOM	2384	CD2	TYR	300	24.490	53.640	54.827	1.00 31.37	'AAAA
MOTA	23.85	CET	TYR		24.501		53.916	1.00 35.88	
-									
ATOM	2386	CZ	_TYR	300	25.448	54.689	52.913	1.00 38.44	AAAA
ATOM	2387	OH	TYR	300	25.417	55.700	51.990	1.00 33.41	AAAA
ATOM	2388	С	TYR	300	26.280	49.515	56.921	1.00 22.80	
MOTA	2389	0	TYR	300	26.895	49.643	57.983	1.00 19.14	AAAA
	2390	N	ALA	301	25.374		56.705	1.00 23.08	
ATOM									
ATOM	2391	CA	ALA	301	25.009	47.589	57.719	1.00 21.68	AAAA
MOTA.	2392	CB	ALA	301	23.893	46.687	57.198	1.00 19.52	
MOTA	2393	С	ALA	301	26.216	46.762	58.098	1.00 23.49	AAAA
ATOM	2394	0	ALA	301	26.507	46.570	59.274	1.00 21.21	AAAA
ATOM	2395	N	LEU	302	26.904		57.072		
								1.00 23.19	
ATOM	2396	CA	LEU	302	28.090	45.463	57.234	1.00 20.66	AAAA
MOTA	2397	CB	LEU	. 302	28.602	45.057	55.844	1.00 23.31	AAAA
ATOM	2398	CG	LEU	302	29.932		55.611	1.00 36.66	AAAA
ATOM	2399	CD1	LEU	302	29.979	43.849	54.170	1.00 38.41	AAAA
ATOM	2400		LEU	302	31.104		55.879	1.00 28.52	
MOTA	2401	С	LEU	302	29.165		58.012	1.00 22.08	
ATOM	2402	0	LEU	302	29.653	45.713	59.020	1.00 20.43	AAAA
ATOM	2403	N	ALA	303	29.517	47.401	57.549	1.00 19.58	
MOTA	2404	CA	ALA	303	30.567		58.197	1.00 19.77	
ATOM	2405	CB	ALA	303	30.816	49.460	57.432	1.00 21.69	AAAA
ATOM	2406	С	ALA	303	30.324	48.485	59.657	1.00 19.19	AAAA
				303					
ATOM	2407	0	ALA		31.216		60.489	1.00 22.51	
ATOM	2408	N	ARG	304	29.128	48.954	59.993	1.00 20.12	AAAA
ATOM	2409	CA	ARG	304	28.872	49.296	61.377	1.00 18.04	AAAA
ATOM	2410	CB	ARG	304 .	27.566		61.511	1.00 21.09	
ATOM	2411	CG	ARG	304	27.532	51.481	60.792	1.00 24.34	AAAA
ATOM	2412	CD	ARG	304	26.259	52.259	61.206	1.00 27.09	AAAA
ATOM	2413	NE	ARG	304	25.090	51.398	61.116	1.00 45.73	AAAA
ATOM	2414	CZ	ARG	304	23.965	51.549	61.808	1.00 39.82	
ATOM	2415	NH1	ARG	304	23.813	52.550	62.677	1.00 28.40	AAAA
ATOM	2416	NH2	ARG	304	22.991	50.667	61.647	1.00 41.77	AAAA
ATOM	2417	С	ARG	304	28.794	48.073	62.280	1.00 21.00	AAAA
ATOM	2418	0	ARG	304	29.313	48.087	63.397	1.00 19.45	AAAA
ATOM	2419	N	ALA	305	28.159	47.008	61.796	1.00 19.93	AAAA
									•
MOTA	2420	CA	ALA	305	28.002	45.809	62.610	1.00 18.70	AAAA
MOTA	2421	CB	ALA	305	26.998	44.830	61.933	1.00 18.26	· AAAA
ATOM	2422	С	ALA	305	29.311	45.109	62.915	1.00 16.46	AAAA
MOTA	2423	0	ALA	305	29.564	44.736	64.061	1.00 19.49	AAAA .
MOTA	2424	N	TRP	306	30.152	44.909	61.905	1.00 `1.92	AAAA
ATOM	2425	CA	TRP	306	31.423	44.268	62.183	1.00 18.99	AAAA
ATOM	2426	CB	TRP	306	32.151	43.865	60.902	1.00 17.96	AAAA
MOTA	2427	CG	TRP	306	31.632	42.564	60.333	1.00 21.34	AAAA
	2428	CD2		306	31.852	42.058	59.014		AAAA
ATOM								1.00 16.55	
MOTA	2429	CE2	TRP	306	31.243	40.785	58.949	1.00 19.37	AAAA
ATOM	2430	CE3	TRP	306	32.507	42.556	57.878	1.00 -17.80	AAAA
ATOM .		CD1		306	30.919	41.610	60.995	1.00 19.88	AAAA
MOTA	2432	NE1		306	30.680	40.535	60.170	1.00 15.95	AAAA
MOTA	2433	CZ2		306	31.270	40.002	57.787	1.00 24.85	AAAA
ATOM	2434	CZ3		306	32.534	41.781	56.725	1.00 29.69	AAAA
MOTA	2435	CH2	TRP	306 ·	31.917	40.513	56.691	1.00 17.04	AAAA
MOTA	2436	С	TRP	306	32.289	45.168	63.018	1.00 20.26	AAAA
ATOM	2437	Ó	TRP	306	33.159	44.726	63.752	1.00 21.20	AAAA
							62.911		
ATOM	2438	N	THR	307	32.061	46.491		1.00 18.60	AAAA
ATOM	2439	ÇA	THR	307	32.843	47.412	63.722	1.00 16.88	AAAA
MOTA	2440	CB	THR	307	32.579	48.885	63.312	1.00 22.05	AAAA
		oG1		307		49.132	62.051	1.00 21.58	AAAA
MOTA	2441				33.218				
ATOM	3442	CG2.	THR	307	33.126	49.857	64.356	1.00 24.86	AAAA

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ATOM	2443	С	THR	307	32.493	47.146	65.187	1.00 17.47	AAAA
	2444	0	THR	307	33.377	47.142	66.039	1.00 18.94	AAAA
ATOM				308	31.216	46.901	65.487	1.00 19.97	AAAA
ATOM	2445	N	LEU						
ATOM	2446	CA	LEU	308	30.834	46.587	66.866	1.00 22.54	AAAA
	2447	CB	LEU	308	29.318	46.365	66.989	1.00 21.13	AAAA
MOTA					28.415	47.579	66.751	1.00 22.82	AAAA
ATOM	2448	CG	LEU	308					
ATOM	2449	CD1	LEU	308	26.937	47.219	67.023	1.00 25.01	AAAA
			LEU	308	28.870	48.710	67.685	1.00 29.09	AAAA
MOTA	2450						67.336	1.00 22.98	AAAA
ATOM	2451	С	LEU	308	31.578	45.331			
ATOM	2452	. 0	LEU	308	32.056	45.250	68.479	1.00 22.27	AAAA
		N	ILE	309	31.677	44.342	66.454	1.00 22.54	AAAA
MOTA	2453						66.801	1.00 17.09	AAAA
MOTA	2454	CA	ILE	309	32.377	43.114			
MOTA	2455	CB	ILE	309	32.318	42.073	65.664	1.00 18.12	AAAA ·
	2456		ILE	309	33.170	40.870	66.033	1.00 24.16	AAAA
ATOM						41.655	65.399	1.00 18.26	AAAA
ATOM	2457		ILE	309	30.871				
ATOM	2458	CD1	ILE	309	30.205	40.989	66.586	1.00 26.57	AAAA
		C	ILE	309	33.849	43.410	67.067	1.00 20.84	AAAA
MOTA	2459						68.031	1.00 25.20	AAAA
ATOM	2460	0	ILE	309	34.426	42.905			
ATOM	2461	N	, TRP	310	34.466	44.223	66.214	1.00 16.86	AAAA
		.CA	TRP	310	35.888	44.517	66.411	1.00 17.86	AAAA
MOTA	2462					45.319	65.235	1.00 14.83	AAAA
MOTA	2463	CB	TRP	310	36.439				
MOTA	2464	CG	TRP	310	37.879	45.648	65.397	1.00 16.63	AAAA
	2465	CD2		310	38.967	44.718	65.560	1.00 18.62	AAAA
MOTA					40.131	45.478	65.799	1.00 25.60	AAAA
MOTA	2466	CE2	TRP	310					
ATOM	2467	CE3	TRP	310	39.069	43.319	65.529	1.00 24.06	AAAA
	2468	CD1	TRP	310	38.418	46.895	65.533	1.00 19.82	AAAA
ATOM					39.768	46.801	65.777	1.00 25.84	AAAA
ATOM	2469	NE1		310					AAAA
MOTA	2470	CZ2	TRP	310	41.383	44.887	66.006	1.00 26.14	
ATOM	2471	CZ3	TRP	310	40.308	42.730	65.735	1.00 24.89	AAAA
			TRP	310	41.452	43.515	65.971	1.00 24.96	AAAA
MOTA	2472	CH2						1.00 20.86	AAAA
ATOM	2473	С	TRP	310	36.112	45.263	67.733		
ATOM	2474	0	TRP	310	37.050	44.957	68.476	1.00 21.38	AAAA
			CYS	311	35.242	46.226	68.030	1.00 24.22	AAAA
MOTA	2475	N					69.280	1.00 27.66	AAAA
ATOM	2476	CA	CYS	311	35.349	46.971			
ATOM	2477	CB	CYS	311	34.297	48.097	69.343	1.00 25.37	AAAA
	2478	SG	CYS	311	34.618	49.528	68.253	1.00 27.22	AAAA
ATOM	_						70.490	1.00 22.95	AAAA
ATOM	2479	С	CYS	311	35.224				AAAA
MOTA	2480	0	CYS	311	35.986	46.180	71.441	1.00 25.47	
	2481	N	GLU	312	34.284	45.089	70.457	1.00 17.03	AAAA
ATOM					34.120	44.129	71.569	1.00 22.44	AAAA
ATOM	2482	CA	GLU	312					AAAA
ATOM	2483	CB	GLU	312	33.011	43.110	71.280	1.00 20.81	
ATOM	2484	CG	GLU	312	31.856	43.048	72.258	1.00 43.65	AAAA
		CD	GLU	312	32,265	42.971	73.717	1.00 29.63	AAAA
ATOM	2485				_		74.119	1.00 38.85	AAAA
ATOM	2486	OEl	GLU	312	33.022	42.059			
ATOM	2487	OE2	GLU	312	31.804	43.844	74.473	1.00 53.22	AAAA
		С	GLI	312	35.395	43.309	71.778	1.00 27.47	AAAA
MOTA	2488				35.899	43.178	72:895	1.00 22.33	AAAA
MOTA	2489	0	GL\.	312	_				AAAA
ATOM	2490	N	LEC	313	35.899	42.723	70.696	1.00 23.82	
MOTA	2491	CA	LEU	313	37.101	41.889	70.771	1.00 20.72	AAAA
					37.380	41.222	69.422	1.00 27.82	AAAA
ATOM	2492	CB	LEU	313				1.00 33.55	AAAA
ATOM	2493	CG	LEU	313	36.403	40.167	68.903		
	2494	CD1	LEU	313	36.839	39.738	67.512	. 1.00 24.03	AAAA
ATOM		CD2		313	36.379	38.981	69.846	1.00 28.20	AAAA
MOTA	2495	CDZ						1.00 18.21	AAAA
ATOM	2496	С	LEU	313	38.343	42.670	71.181		
ATOM	2497	0	LEU	313	39.119	42.205	72.017	1.00 21.48	AAAA
				314	38.492	43.848	70.580	1.00 19.41	AAAA
MOTA	2498	N	SER					1.00 28.26	AAAA
ATOM	2499	CA	SER	314	39.627	44.753	70.775		
ATOM	2500	CB	SER	314 ·	39.625	45.821	69.663	1.00 22.55	AAAA
				314	40.732	46.696	69.759	1.00 61.92	AAAA
ATOM	2501	OG	SER					1.00 30.18	AAAA
MOTA	2502	С	SER	314	39.619	45.429	72.144	1.00 50.20	
ATOM	2503	0	SER	314	40.631	45.969	72.590	1.00 25.04	AAAA
		N	GLY	315	38.477	45.407	72.806	1.00 28.04	AAAA
ATOM	2504				38.393		74.119	1.00 33.84	AAAA
MOTA	2505	CA	GLY	315		46.009		1.00 35.01	AAAA
MOTA	2506	С	GLY	315	38.324	47.518	74.105	1.00 36.93	
	2507	ō	GLY	315	38.811	48.178	75.022	1.00 37.00	AAAA
MOTA					37.739	48.090	73.065	1.00 31.33	. AAAA
ATOM	2508	N	ARG	316	31.133	30.070			

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MOTA	2509	CA	ARG	316	37.631	49.536	73.042	1.00 39.10	AAAA
ATOM	2510	CB	ARG	316	38.347	50.108	71.830	1.00 45.15	AAAA
MOTA	2511	CG	ARG		37.722	49.834		1.00 46.02	AAAA
ATOM	2512	CD	ARG	316	38.620	50.459	69.449	1.00 44.83	AAAA
MOTA	2513	NE	ARG	316	39.898	49.767	69.357	1.00 37.91	AAAA
ATOM	2514	CZ	ARG		40.945	50.219		1.00 27.39	
									AAAA
atom	2515	NH1	ARG	316	40.854	51.371	68.034	1.00 50.24	AAAA
MOTA	2516	NH2	ARG	316	42.054	49.493	68.572	1.00 34.51	AAAA
MOTA	2517	С	ARG	316	36.179	49.984		1.00 35.43	AAAA ·
ATOM	2518	0	ARG	316	35.292	49.271		1.00 30.71	AAAA
ATOM	2519	N	GLU	317	35.931	51.162	73.612	1.00 34.06	AAAA
ATOM	2520	CA	GLU	317	34.569	51.663	73.671	1.00 37.96	AAAA
ATOM	2521	CB	GLU	317	34.481	52.914		1.00 43.60	
									AAAA
ATOM	2522	CG	GLU		33.961	52.630		1.00 60.36	AAAA
ATOM	2523	CD	GLU	317	34.768	51.575	76.701	1.00 70.70	AAAA
ATOM	2524	OE1	GLU	317	34.375	51.217	77.832	1.00 76.71	AAAA
	2525	OE2		317	35.793	51.104			
MOTA								1.00 78.36	AAAA
MOTA	2526	С	GLU	317	34.068	51.958	72.280	1.00 35.65	AAAA
ATOM	2527	0	GLU	317	34.843	52.322	71.390	1.00 32.91	AAAA
ATOM	2528	N	VAL	318	32.767	51.772	72.094	1.00 30.52	AAAA
ATOM	2529	CA	VAL	318	32.138	52.012	70.808	1.00 37.04	AAAA
MOTA	2530	CB	VAL	318	30.877	51.138	70.638	1.00 36.48	AAAA
MOTA	2531	CG1	VAL	318	30.278	51.366	69 ⁻ . 268	1.00 40.43	AAAA
ATOM	2532		VAL	318	31.222	49.674	70.846	1.00 33.75	
									AAAA
MOTA	2533	С	VAL	318	31.719	53.465	70.737	1.00 28.96	AAAA
ATOM	2534	0	VAL	318	30.930	53.915	71.556	1.00 33.56	AAAA
ATOM	2535	N	PRO	319	32.258	54.229	69.773	1.00 29.20	AAAA
ATOM	2536	CD	PRO	319	33.243	53.924	68.726	1.00 31.62	
									AAAA
ATOM	2537	CA	PRO	319	31.858	55.637	69.684	1.00 28.99	AAAA
ATOM	2538	CB	PRO	319	32.709	56.154	68.528	1.00 32.17	AAAA
ATOM	2539	CG	PRO	319	32.850	54.926	67.664	1.00 41.36	AAAA
ATOM	2540		PRO	319	30.365	55.680	69.377	1.00 36.95	
		C							AAAA
ATOM	2541	0	PRO	319	29.847	54.795	68.695	1.00 32.86	AAAA
ATOM	2542	N	GLU	320	29.646	56.683	69.855	1.00 34.61	AAAA
ATOM	2543	CA	GLU	320	28.230	56.657	69.544	1.00 35.13	AAAA
ATOM	2544	CB	GLU	320	27.419	57.416	70.595	1.00 52.97	AAAA
ATOM	2545	CG	GLU	320	27.751	58.875	70.738	1.00 56.06	AAAA
ATOM	2546	CD	GLU	320	26.822	59.558	71.721	1.00 65.58	AAAA
ATOM	2547		GLU	320	25.604	59.619	71.444	1.00 64.27	AAAA
MOTA	2548		GLU	320	27.306	60.022	72.775	1.00 72.99	AAAA
MOTA	2549	С	GLU	320	27.943	57.192	68.153	1.00 35.13	AAAA
MOTA	2550	0	GLU	320	26.916	56.879	67.565	1.00 37.43	AAAA
ATOM	2551	N	LYS	321	28.880	57.953	67.604	1.00 28.22	AAAA
ATOM	2552	CA	LYS	321	28.700	58.555	66.289	1.00 36.58	AAAA
ATOM	2553	CB	LYS	321	28.666	60.071	66.454	1.00 44.87	AAAA
ATOM	2554	CG	LYS	321	29.987	60.606	67.023	1.00 55.73	· `AAA
ATOM	2555	CD	LYS	321	30.305	60.020	68.410	1.00 57.27	LAAA
ATOM	2556	CE	LYS	321	31.733	60.310	68.840	1.00 54.59	AAA.
MOTA	2557	NZ	LYS	321	32.024	61.774	68.848	1.00 67.47	AAAA
ATOM	2558	С	LYS	321	29.823	58.211	65.315	1.00 34.44	AAAA
ATOM	2559	0	LYS	321	30.912	57.818	65.731	1.00 33.83	AAAA
						_			
ATOM	2560	N	LEU	322	29.549	58.354	64.019	1.00 30.21	AAAA
MOTA	2561	CA	LEU	322	30.575	58.135	62.998	1.00 29.45	AAAA
ATOM	2562	CB	LEU	322	29.966	57.677	61.677	1.00 32.21	AAAA
	2563		LEU		29.240	56.338	61.651	1.00 38.94	
MOTA				322					AAAA
MOTA	2564	CD1	LEU	322	29.008	55.977	60.186	1.00 38.44	AAAA
ATOM	2565	CD2	LEU	322	30.072	55.261	62.337	1.00 42.11	AAAA
MOTA	2566		LEU	322	31.228	59.503	62.783	1.00 33.28	AAAA
ATOM	2567		LEU	322	30.544	60.519	62.872	1.00 31.45	AAAA
MOTA	2568	N	ASN	323	32.533	59.539	62.519	1.00 34.38	AAAA
ATOM	2569		ASN	323	33.208	60.824	62.294	1.00 36.53	AAAA
	2570		ASN	323	34.701	60.737	62.600	1.00 42.85	AAAA
ATOM									
ATOM	2571		ASN	323	35.484	60.081	61.480	1.00 50.51	AAAA
ATOM ·	2572	OD1	ASN	323	35.215	58.942	61.109	1.00 51.23	AAAA
ATOM	2573	ND2		323	36.455	60.807	60.928	1.00 60.23	AAAA
					33.027	61.171	60.822		
MOTA	2574	.C	asn	323	33.041	AT.T/T	20.022	1.00 34.69	AAAA

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3001	2575	0	ASN	323	32.429	60.395	60.075	1.00 34.06	AAAA
MOTA							60.390	1.00 23.73	AAAA
ATOM	2576	N	ASN	324	33.551	62.317			
ATOM	2577	CA	ASN	324	33.385	62.720	58.995	1.00 31.06	AAAA
ATOM	2578	CB	ASN	324	33.868	64.155	58.784	1.00 36.07	AAAA
			ASN	324	32.974	65.163	59.450	1.00 45.77	AAAA
MOTA	2579	CG							
MOTA	2580	OD1	ASN	324	31.765	65 ⁻ . 189	59.206	1.00 39.55	AAAA
ATOM	2581	ND2	ASN	324	33.555	66.008	60.291	1.00 39.12	AAAA
		c	ASN	324	34.047	61.810	57.971	1.00 24.87	AAAA
ATOM	2582							1.00 31.91	AAAA
ATOM	2583	0	ASN	324	33.451	61.483	56.941		
MOTA	2584	N	LYS	325	35.276	61.405	58.250	1.00 27.73	AAAA
MOTA	2585	CA	LYS	325	- 35.991	60.538	57.333	1.00 29 <i>.</i> 55	AAAA
					37.351	60.182	57.929	1.00 37.43	AAAA
MOTA	2586	CB	LYS	325					
MOTA	2587	CG	LYS	325	38.250	59.396	57.004	1.00 44.84	AAAA
MOTA	2588	CD	LYS	325	39.684	59.435	57.502	1.00 50.89	AAAA
		CE	LYS	325	40.191	60.873	57.561	1.00 54.82	AAAA
MOTA	2589					60.980	57.969	1.00 65.70	AAAA
ATOM	2590	NZ	LYS	325	41.621				
ATOM	2591	С	LYS	325	35.161	59.279	57.078	1.00 27.99	AAAA
ATOM	2592	0	LYS	325	35.016	58.836	55.938	1.00 31.80	AAAA
				326	34.602	58.721	58.142	1.00 26.07	AAAA
ATOM	2593	· N	ALA						
ATOM	2594	CA	ALA	326	33.781	57.506	58.030	1.00 24.38	AAAA
MOTA	2595	CB	ALA	326	33.470	56.982	59.428	1.00 27.34	AAAA
	2596	c	ALA	326	32.478	57.709	57.231	1.00 25.78	AAAA
MOTA						56.890	56.369	1.00 27.37	AAAA
ATOM	2597	0	ALA	326	32.131				
ATOM	2598	N	LYS	327	31.749	58.790	57.496	1.00 27.31	AAAA
ATOM	2599	CA	LYS	327	30.502	59.027	56.758	1.00 28.56	AAAA
	2600	CB	LYS	327	29.759	60.251	57.313	1.00 28.87	AAAA
MOTA					29.491	60.209	58.812	1.00 36.72	AAAA
MOTA	2601	CG	LYS	327					
ATOM	2602	CD	LYS	327	· 28.643	61.407	59.255	1.00 40.34	, AAAA
ATOM	2603	CE	LYS	327	28.645	61.594	60.769	1.00 38.91	AAAA
		NZ	LYS	327	28.163	60.429	61.556	1.00 47.67	AAAA
ATOM-	2604					59.244	55.269	1.00 29.15	AAAA
MOTA	2605	С	LYS		30.792				
MOTA	2606	0	LYS	327	30.097	58.719	54.393	1.00 27.76	AAAA
ATOM	2607	N	GLU	328	31.829	60.015	54.972	1.00 31.59	AAAA
			GLU	328	32.167	60.265	53.581	1.00 28.93	AAAA
ATOM	2608	CA						1.00 32.30	AAAA
MOTA	2609	CB	GLU	328	33.257	61.332	53.515		
ATOM	2610	CG	GLU	328	32.745	62.652	54.067	1.00 47.50	AAAA
ATOM	2611	CD	GLU	328	33.764	63.772	54.032	1.00 46.67	AAAA
				328	34.325	64.037	52.951	1.00 56.88	AAAA
MOTA	2612	OE1	GLU						AAAA
MOTA	2613	OE2	GLU	328	33.984	64.402	55.087	1.00 42.24	
ATOM	.2614	С	GLU	328	32.575	58.975	52.871	1.00 30.46	AAAA
MOTA	2615	0	GLU	328	32.226	58.753	51.704	1.00 26.29	AAAA
					33.292	58.112	53.584	1.00 24.93	AAAA
MOTA	2616	N	LEU	329					AAAA
MOTA	2617	CA	LEU	329	33.701	56.828	53.017	1.00 24.80	
MOTA	2618	CB	LEU	329	34.478	56.003	54.053	1.00 25 <i>.</i> 70	AAAA
	2619	CG	LEU	329	34.730	54.522	53.703	1.00 19.71	AAAA
MOTA					5.569	54.413	52.430	1.00 25.26	AAAA
MOTA	2620		LEU	329					AAAA
MOTA	2621	CD2	LEU	329	55.412	53.833	54.863	1.00 24.73	
ATOM	2622	С	LEU	329	2.443 د	56.059	52.603	1.00 23.50	AAAA
	2623	ō	LEU	329	32.310	55.650	51.453	1.00 25.60	AAAA
MOTA							53.539	1.00 23.02	AAAA
ATOM	2624	N	LEU	330	31.516	55.881			
ATOM	2625	CA	LEU	330	30.289	55.145	53.242	1.00 23.85	AAAA
ATOM	2626	CB	LEU	330	29.414	55.030	54.484	1.00 21.74	AAAA
			LEU	330	30.039	54.252	55.642	1.00 25.29	AAAA
ATOM	2627	CG						1.00 30.58	. AAAA
ATOM	2628	CD1	LEU	330	28.984	54.053	56.724		
ATOM	2629	CD2	LEU	330	30.538	52.905	55.168	1.00 22.44	AAAA
	2630	C	LEU	330	29.491	55.769	52.113	1.00 26.94	AAAA
ATOM					28.968	55.060	51.252	1.00 26.65	AAAA
MOTA	2631	Ö	LEU	330					
ATOM	2632	N	LYS	331	29.404	57.097	52.111	1.00 30.82	AAAA
ATOM	2633	CA	LYS	331	28.667	57.795	51.066	1.00 29.53	AAAA
			LYS	331	28.537	59.292	51.407	1.00 29.67	AAAA
ATOM	2634	CB						1.00 36.06	AAAA
MOTA	2635	CG	LYS	331	27.814	59.567	52.714	1.00 30.00	
ATOM	2636	CD	LYS	331	27.688	61.055	52.990	1.00 42.75	AAAA
	2637	CE	LYS	331	26.828	61.737	51.939	1.00 53.98	AAAA
MOTA				331	26.634	63.185	52.234	1.00 67.57	AAAA
MOTA	2638	NZ	LYS				40 600	1.00 30.07	AAAA
MOTA	2639	C .	LYS	331	29.315	57.628	49.692	1.00 30.07	
ATOM	2640		LYS	331	28.634	57.759	48.672	1.00 36.20	AAAA

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ATOM	2641	N	SER	332	30.608	57.305	49.657	1.00 30.08	AAAA
ATOM	2642	CA	SER		31.322		48.385	1.00 33.35	AAAA
MOTA	2643	CB	SER		32.834		48.590	1.00 40.36	AAAA
ATOM	2644	OG	SER	. 332	33.396	56.169	49.219	1.00 34.04	AAAA
MOTA	2645	С	SER	332	31.061	55.821	47.693	1.00 37.72	AAAA
ATOM	2646	0	SER		31.354		46.507	1.00 30.78	AAAA
ATOM	2647	N	ILE	333	30.521	54.865	48.440	1.00 30.61	AAAA
MOTA	2648	CA	ILE	333	30.219	53.547	47.899	1.00 37.59	AAAA
	-								
MOTA	2649	CB	ILE		29.901	52.551	49.022	1.00 33.59	AAAA
ATOM	2650	CG2	LILE	333	29.738	51.146	48.442	1.00 37.05	AAAA
ATOM	2651	CG1	ILE	333	31.015	52.564	50.065	1.00 38.95	AAAA
							51.282		
MOTA	2652		ILE		30.706	51.727		1.00 46.83	AAAA
MOTA	2653	C	ILE	333	28.990	53.620	46 ⁻ .998	1.00 43.41	AAAA
ATOM	2654	0	ILE	333	27.889	53.876	47.479	1.00 46,24	AAAA
	2655	N	ASP	334	29.158	53.423	45.696	1.00 47.97	
ATOM									AAAA
ATOM	2656	CA	ASP	334	27.976	53.447	44.847	1.00 53.47	- AAAA
ATOM	2657	CB	ASP	334	28.333	53.535	43.358	1.00 61.52	AAAA
ATOM	2658	CG	ASP	334	29.223	52.406	42.897	1.00 64.75	AAAA
ATOM	2659		ASP	334	29.379	52.248	41.666	1.00 66.93	AAAA
'ATOM	2660	OD2	ASP	334	29.779	51.691	43.758	1.00 65.93 -	AAAA
ATOM	2661	С	ASP	334	27.248	52.144	45.161	1.00 51.83	AAAA
				334					
ATOM	2662	0	ASP	_	27.626	51.067	44.699	1.00 46.80	AAAA
ATOM	2663	N	PHE	335	26.215	52.249	45.986	1.00 54.96	AAAA
ATOM	2664	CA	PHE	335	25.455	51.080	46.392	1.00 50.60	AAAA
ATOM	2665	CB	PHE	335	25.413	51.003	47.920	1.00 39.55	AAAA
MOTA	2666	CG	PHE	335	24.380	50.054	48.440	1.00 37.98	AAAA
ATOM	2667	CD1	PHE	335	24.389	48.715	48.054	1.00 46.72	AAAA
ATOM	2668	CD2	PHE	335	23.362	50.506	49.262	1.00 34.23	AAAA
	2669		PHE	335	23.389	47.842	48.478	1.00 49.80	
ATOM									AAAA
ATOM	2670	CE2	PHE	335	22.361	49.644	49.689	1.00 48.51	AAAA
ATOM	2671	CZ	PHE	335	22.373	48.309	49.296	1.00 40.44	AAAA
MOTA	2672	C	PHE	335	24.033	51.000	45.839	1.00 54.52	AAAA
MOTA	2673	0	PHE	335	23.603	49.939	45.379	1.00 59.24	AAAA
ATOM	2674	N	GLU	336	23.302	52.108	45.888	1.00 50.94	AAAA
ATOM	2675	CA	GLU	336	21.923	52.119	45.406	1.00 57.05	AAAA
ATOM	2676	CB	GLU	336	21.853	51.751	43.924	1.00 60.27	AAAA
ATOM	2677	CG	GLU	336	20.430	51.627	43.422	1.00 68.55	AAAA
ATOM	2678	CD	GLU	336	20.352	51.126	42.001	1.00 80.03	AAAA
ATOM	2679	OF1	GLU	336	20.860	50.013	41.735	1.00 84.64	AAAA
ATÓM	2680		GLU	336	19.777	51.841	41.153	1.00 80.68	AAAA
MOTA	2681	С	GLU	336	21.065	51.135	46.201	1.00 55.73	AAAA
ATOM	2682	0	GLU	336	21.219	49.917	46.089	1.00 51.33	AAAA
MOTA	2683	N	GLU	337	20.151	51.679	46.992	1.00 49.54	AAAA
				337	19.267		47.821	1.00 48.19	
MOTA	2684	CA	GLU			50.880			AAAA
ATOM	2685	CB	GLU	337	18.510	51.822	48.764	1.00 47.73	AAAA
MOTA	2686	CG	GLU	337	18.084	51.205	50.077	1.00 55.69	AAAA
ATOM	2687	CD	GLU	337	19.269	50.720	50.904	1.00 50.17	AAAA
				337	20.111	51.548	51.345	1.00 36.03	
atom	2688	OE1							AAAA
ATOM	2689	OE2	GLU	337	19.358	49.494	51.105	1.00 51.25	AAAA
ATOM	2690	С	GLU	337	18.294	50.083	46.936	1.00 49.13	AAAA
	2691	0	GLU	337	17.816	50.588	45.916	1.00 48.61	AAAA
ATOM									
MOTA	2692	N	PHE	338	18.015	48.837	47.313	1.00 48.15	AAAA
ATOM	2693	CA	PHE	338	17.092	48.000	46.547	1.00 48.12	AAAA
MOTA	2694	CB	PHE	338	16.870	46.658	47.249	1.00.54.54	AAAA
MOTA	2695	CG	PHE	338	15.883	45.777	46.548	1.00 57.22	AAAA
ATOM .	2696	CD1	PHE	338	16.115	45.366	45.243	1.00 60.01	AAAA
ATOM	2697	CD2	PHE	338	14.699	45.398	47.171	1.00 55.04	AAAA
	•	CE1		338	15.185	44.597	44.566	1.00 60.84	AAAA
MOTA	2698								
ATOM	2699	CE2	PHE	338	13.758	44.624	46.497	1.00 59.41	AAAA
MOTA	2700	CZ	PHE	338	14.002	44.224	45.189	1.00 57.18	AAAA
ATOM	2701	c	PHE	338	15.755	48.714	46.380	1.00 45.46	AAAA
MOTA	2702	0	PHE	338	15.274	48.900	45.263	1.00 51.11	AAAA
ATOM	2703	N	ASP	339	15.154	49.098	47.501	1.00 40.38	AAAA
ATOM ·	2704	CA	ASP	339	13.890	49.820.	47.488	1.00 49.97	AAAA
			ASP	339	13.270	49.821	48.886	1.00 53.23	AAAA
MOTA	2705			•				1.00 55.25	
ATOM	2706	CG	ASP	339	12.000	50.659	48.968	1.00 57.40	AAAA
		•		•		_		•	•

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ATOM	2707	ODI	ASP	339	12.039	51.858	48.616	1.00 53.79	AAAA
	2707		ASP	339	10.963	50.118	49.401	1.00 51.15	AAAA
MOTA					14.215	51.248	47.076	1.00 55.06	AAAA
ATOM	2709	C	ASP	339		51.922			
MOTA	2710	0	ASP	339	14.994		47.748	1.00 56.47	AAAA
MOTA	2711	N	ASP	340	13.623	51.708	45.978	1.00 58.46	AAAA
MOTA	2712	CA	ASP	340	13.874	53.059	45.484	1.00 67.72	AAAA
MOTA	2713	CB	ASP	340	12.683	53.559	44.664	1.00 71.52	AAAA
MOTA	2714	CG	ASP	340	12.611	52.913	43.295	1.00 79.72	AAAA
ATOM	2715	OD1	ASP	340	12.528	51.667	43.224	1.00 86.74	AAAA
ATOM	2716	OD2	ASP	340	12.640	53.655	42.288	1.00 83.40	AAAA
ATOM	2717	С	ASP	340	14.209	54.072	46.572	1.00 69.65	AAAA
ATOM	2718	0	ASP	340	15.204	54.794	46.463	1.00 70.13	AAAA
ATOM	2719	N	GLU	341	13.392	54.130	47.620	1.00 67.11	AAAA
ATOM	2720	CA	GLU	341	13.668	55.077	48.689	1.00 67.87	AAAA
MOTA	2721	CB	GLU	341	13.195	56.478	48.278	1.00 74.87	AAAA
MOTA	2722	CG	GLU	341	13.502	57.576	49.298	1.00 82.72	AAAA
	2723	CD	GLU	341	13.162	58.974	48.790	1.00 90.80	AAAA
MOTA					11.988	59.215	48.431	1.00 90.38	AAAA
ATOM	2724		GLU	341		59.835	48.752	1.00 93.36	AAAA
MOTA	2725		GLU	341	14.072				
MOTA	2726	C	GLU	341	13.101	54.719	50.058	1.00 60.22	AAAA
MOTA	2727	0	GLU	341	11.929	54.955	50.347	1.00 58.81	AAAA
ATOM	2728	N	VAL	342	13.956	54.144	50.897	1.00 57.28	AAAA
ATOM	2729	CA	VAL	342	13.594	53.781	52.262	1.00 52.09	AAAA
MOTA	2730	CB	VAL	342	14.195	52.419	52.669	1.00 53.17	AAAA
MOTA	2731	CG1	VAL	342	13.730	52.042	54.070	1.00 46.16	AAAA
ATOM	2732	CG2	VAL	342	13.815	51.356	51.663	1.00 59.09	AAAA
ATOM	2733	С	VAL	342	14.263	54.843	53.124	1.00 53.31	AAAA
ATOM	2734	0	VAL	342	13.763	55.230	54.185	1.00 57.79	AAAA
ATOM	2735	N	ASP	343	15.398	55.306	52.610	1.00 46.24	AAAA
ATOM	2736	CA	ASP	343	16.268	56.289	53.243	1.00 42.60	AAAA
ATOM	2737	CB	ASP	343	15.521	57.510	53.781	1.00 43.88	AAAA
ATOM	2738	CG	ASP	343	16.480	58.581	54.290	1.00 46.82	AAAA
MOTA	2739		ASP	343	16.028	59.581	54.887	1.00 46.16	AAAA
	2740		ASP	343 .	17.700	58.414	54.075	1.00 33.01	AAAA
MOTA	2741	C	ASP	343	17.012	55.636	54.395	1.00 35.45	AAAA
ATOM	•			343	16.487	55.480	55.502	1.00 29.39	AAAA
ATOM	2742	0	ASP		18.247	55.249	54.124	1.00 30.51	AAAA
ATOM	2743	N	ARG	344			55.140	1.00 29.43	AAAA
ATOM	2744	CA	ARG	344	19.059	54.613		1.00 30.10	AAAA
MOTA	2745	СВ	ARG	344	19.736	53.377	54.561		AAAA
MOTA	2746	CG	ARG	344	18.803	52.258	54.180	1.00 33.95	AAAA
MOTA	2747	CD	ARG	344	17.981	51.770	55.365	1.00 20.92	
ATOM	2748	ΝE	ARG	344	17.120	50.673	54.936	1.00 29.72	AAAA
ATOM	2749	cz	ARG	344	16.110	50.176	55.639	1.00 29.13	AAAA
ATOM	2750	NH1		344	15.805	50.668	56.835	1.00 29.63	AAAA
MOTA	2751	NH2	ARG	344	15.379	49.198		1.00 27.19	AAAA.
ATOM	2752	С	ARG	344	20.116	55.769	55.660	1.00 34.31	AAAA
MOTA	2753	0	ARG	344	21.005	5557	56.391	1.00 29.09	AAAA
ATOM	2754	N	SER	345	20.011	56. 45	55.294	1.00 28.34	AAAA
MOTA	2755	CA	SER	345	20.999	57.839	55.715	1.00 30.95	AAAA
ATOM	2756	CB	SER	345	20.669	59.199	55.109	1.00 29.56	AAAA
ATOM	2757	OG	SER	345	19.429	59.648	55.610	1.00 29.38	AAAA
ATOM	2758	C	SER	345	21.137	57.988	57.230	1.00 30.92	AAAA
ATOM	2759	ō	SER	345	22.155	58.488	57.718	1.00 31.15	AAAA
MOTA	2760	N	TYR	346	20.116	57.576	57.975	1.00 25.64	AAAA
	2761	CA	TYR	346	20.158	57.659	59.433	1.00 26.81	AAAA
ATOM					18.823	57.189	60.006	1.00 34.41	AAAA
MOTA	2762	CB	TYR	346			59.716	1.00 27.35	AAAA
MOTA	2763	CG	TYR	346	18.529	55.723	60.556	1.00 24.87	AAAA
MOTA	2764		TYR	346	19.003	54.708		1.00 24.87	AAAA
MOTA	2765		TYR	346	18.744	53.352	60.278		AAAA
ATOM	2766		TYR	346	17.795	55.358	58.588	1.00 27.70	
ATOM	2767	CE2		346	17.533	54.008	58.297	1.00 26.59	AAAA
ATOM	2768	CZ	TYR	346	18.008	53.015	59.145	1.00 33.75	AAAA
ATOM	2769	ОН	TYR	346	17.737	51.691	58.855	1.00 26.06	AAAA
ATOM ·	2770	C	TYR	346	21.277	56.766	59.977	1.00 25.57	AAAA
ATOM	2771	0	TYR	346	21.769	56.970	61.085	1.00 28.07	AAAA
ATOM	2772	N .	MET	347	21.666	55.761	59.198	1.00 29.08	AAAA
ALON				•	•		-		•

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ATOM	2773	CA	MET	347	22.720	54.837	59.622	1.00 24.19	AAAA
	2774	СВ	MET	347	22.844	53.678	58.628	1.00 24.87	AAAA
ATOM	2775	CG	MET	347	21.609	52.806	58.543	1.00 23.66	AAAA
MOTA	2776	SD	MET	347	21.780	51.503	57.267	1.00 27.02	AAAA
ATOM	2777	CE	MET	347	22.115	52.375	55.896	1.00 37.69	AAAA
ATOM	2778	CE	MET	347	24.054	55.540	59.737	1.00 29.45	AAAA
ATOM	2779	ō	MET	347	24.937	55.092	60.479	1.00 28.08	AAAA
MOTA		N	LEU	348	24.188	56.650	59.007	1.00 23.71	AAAA
ATOM	2780	CA	LEU	348	25.418	57.446	58.998	1.00 34.11	AAAA
MOTA	2781		LEU	348	25.463	58.351	57.757	1.00 25.37	AAAA
MOTA	2782	CB CG	LEU	348	25.320	57.785	56.344	1.00 30.38	AAAA
MOTA	2783	CD1		348	25.307	58.944	55.340	1.00 27.44	AAAA
ATOM	2784	CD2		348	26.459	56.814	56.041	1.00 36.44	AAAA
ATOM	2785 2786		LEU	348	25.507	58.332	60.237	1.00 36.09	AAAA
ATOM		С 0	LEU	348	26.561	58.894	60.539	1.00 33.30	AAAA
MOTA	2787	И	GLU	349	24.394	58.445	60.953	1.00 30.51	AAAA
ATOM	2788	CA	GLU	349	24.313	59.292	62.136	1.00 35.53	AAAA
MOTA	2789		GLU	349	22.908	59.896	62.217	1.00 31.35	AAAA
ATOM	2790	CB CG	GLU	349	22.518	60.717	61.006	1.00 29.09	AAAA
MOTA	2791	CD	GLU	349	23.481	61.859	60.746	1.00 31.78	AAAA
ATOM	2792 2793	OE1		349	23.937	62.476	61.730	1.00 30.98	AAAA
MOTA	2794	OE2	GLU	349	23.766	62.155	59.569	1.00 30.67	AAAA
MOTA	2795	C	GLU	349	24.663	58.633	63.471	1.00 38.48	AAAA
MOTA	2796	0	GLU	349	24.727	59.303	64.502	1.00 40.12	AAAA
MOTA	2797	N	THR	350	24.878	57.326	63.461	1.00 33.58	AAAA
ATOM	2798	CA	THR	350	25.221	56.612	64.681	1.00 29.74	AAAA
ATOM ATOM	2799	CB	THR	350	23.992	56.363	65.559	1.00 35.91	AAAA
ATOM	2800	OG1		350	23.421	57.615	65.952	1.00 45.03	AAAA
ATOM	2801	CG2	THR	350	24.382	55.586	66.806	1.00 49.48	AAAA
ATOM	2802	C	THR	350	25.821	55.267	64.330	1.00 30.63	AAAA
ATOM	2803	ō	THR	350	25.535	54.709	63.274	1.00 26.62	AAAA
ATOM	2804	Ň	LEU	351	26.644	54.740	65.225	1.00 29.07	AAAA
ATOM	2805	CA	LEU	351	27.271	53.461	64.972	1.00 24.59	AAAA
ATOM	2806	CB	LEU	351	28.584	53.367	65.757	1.00 29.91	AAAA
ATOM	2807	CG	LEU	351	29.591	52.327	65.267	1.00 39.62	AAAA
MOTA	2808		LEU	351	30.887	52.467	66.039	1.00 37.09	AAAA
ATOM	2809		LEU	351	29.024	50.935	65.415	1.00 54.03	AAAA
ATOM	2810	С	LEU	351	26.314	52.336	65.377	1.00 29.71	AAAA
ATOM	2811	0	LEU	351	26.130	51.364	64.641	1.00 30.53	AAAA
ATOM	2812	N	LYS	352	25.697	52.481	66.543	1.00 28.64	AAAA
ATOM	2813	CA	LYS	352	24.763	51.479	67.061	1.00 32.72	AAAA
ATOM	2814	CB	LYS	352	24.913	51.381	68.581	1.00 27.37	AAAA AAAA
ATOM	2815	CG	LYS	352	26.230	50.787	69.034	1.00 43.48	AAAA
MOTA	2816	CD	LYS	352	26.536	51.068	70.504	1.00 46.77	AAAA
MOTA	2817	CE	LYS	352	25.484	50.538	71.451	1.00 51.52	AAAA
A OM	2818	NZ	LYS	352	25.850	50.859	72.866	1.00 62.08 1.00 32.49	AAAA
A L'OM	2819	С	LYS	352	23.330	51.856	66.731	1.00 32.49	AAAA
A 1 OM	2820	0	LYS	352	22.953	53.010	66.882 66.244	1.00 31.30	AAAA
MOTA	2821	N	ASP	353	22.525	50.916	66.012	1.00 26.50	AAAA
MOTA	2822	CA	ASP	353	21.136	51.286	64.746	1.00 50.09	AAAA
MOTA	2823	CB	ASP	353	20.543	50.635	64.604	1.00 52.79	AAAA
ATOM	2824	CG	ASP	353	20.880	49.176	64.109	1.00 58.55	AAAA
MOTA	2825	OD1	ASP	353	21.980	48.861	64.103	1.00 73.19	AAAA
ATOM	2826		ASP	353	20.040	48.339	67.257	1.00 26.41	AAAA
ATOM	2827	С	ASP	353	20.328	50.930		1.00 25.73	AAAA
ATOM	2828	0	ASP	353	20.806	50.214	68.136 67.385	1.00 30.12	AAAA
MOTA	2829	N	PRO	354	19.118	51.481	66.495	1.00 35.38	AAAA
ATOM	2830	CD	PRO	354	18.428	52.429	68.547	1.00 34.02	AAAA
ATOM	2831	CA	PRO	354	18.276	51.190	68.340	1.00 32.25	AAAA
MOTA	2832	CB	PRO	354	17.091	52.129	66.833	1.00 32.23	AAAA
ATOM	2833	CG	PRO	354	16.974	52.139 49.736	68.512	1.00 34.00	AAAA
ATOM	2834	C	PRO	354	17.838		67.452	1.00 28.28	AAAA
ATOM	2835	0	PRO	354	17.829	49.111 49.190	69.664	1.00 23.89	AAAA
ATOM ·		N	TRP	355	. 17.484		69.669	1.00 33.84	AAAA
ATOM	2837	CA	TRP	355.	17.010	47.818	71.076	1.00 33.84	AAAA
MOTA	2838	CB	TRP	355	16.653	47.363	12.010	1.00 33.04	

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Figure 16-44

MOTA	2839	CG	TRP	355	17.844	46.946.	71.832	1.00 49.97	AAAA
		CD2	TRP	355	18.364	45.622	71.905	1.00 46.60	AAAA
MOTA	2840					45.682	72.639	1.00 54.73	AAAA
MOTA	2841	CE2	TRP	355	19.567				
ATOM	2842	CE3	TRP	355	17.931	44.386	71.419	1.00 46.90	AAAA
	2843		TRP	355	18.723	47.746	72.507	1.00 56.10	AAAA
ATOM						46.991	72.997	1.00 56.07	AAAA
MOTA	2844		TRP	355	19.765				
MOTA	2845	CZ2	TRP	355	20.340	44.552	72.897	1.00 55.25	AAA
ATOM	2846		TRP	355	18.696	43.267	71.674	1.00 50.74	AAAA
						43.356	72.405	1.00 50.68	AAAA
ATOM	2847	CH2	TRP	355	19.887				
ATOM	2848	С	TRP	355	15.789	47.712	68.776	1.00 33.12	AAAA
	2849	ō	TRP	355 .	15.096	48.705	68.550	1.00 29.41	AAAA
MOTA					15.547	46.508	68.263	1.00 23.90	AAAA
ATOM	2850	N	ARG	356					AAAA
MOTA	2851	CA	ARG	356	14.413	46.237	67.387	1.00 23.96	
ATOM	2852	CB	ARG	356	14.892	46.096	6593.5	1.00 22.66	AAAA
				356	15.505	47.385	65.393	1.00 29.06	AAAA
MOTA	2853	CG	ARG		•				AAAA
ATOM	2854	CD	ARG	356	16.291	47.212	64.108	1.00 28.92	
ATOM	2855	NE	ARG	356	16.833	48.503	63.686	1.00 24.73	AAAA
				356	17.733	48.668	62.724	1.00 23.57	AAAA
MOTA	2856	CZ	ARG					1.00 22.15	AAAA
MOTA	2857	NH1	ARG	356	18.209	47.616	62.066		
MOTA	2858	NH2	ARG	356	18.153	49.891	62.418	1.00 22.69	AAAA
		C	ARG	356	13.781	44.944	67.878	1.00 24.89	AAAA
MOTA	2859				13.785	43.925	67.189	1.00 22.25	AAAA
ATOM	2860	0	ARG	356					
ATOM	2861	N	GLY	357	13.231	44.993	69.085	1.00 23.91	AAAA
ATOM	2862	CA	GLY	357	12.631	43.805	69.657	1.00 26.72	AAAA
				357	11.138	43.671	69.465	1.00 26.90	AAAA
MOTA	2863	С	GLY				68.619	1.00 29.87	AAAA
ATOM	2864	0	GLY	357	10.536	44.330			
ATOM	2865	N	GLY	358	10.544	42.797	70.265	1.00 28.22	AAAA
	2866	CA	GLY	358	9.118	42.561	70.188	1.00 30.96	AAAA
ATOM					8.800	41.274	70.920	1.00 30.03	AAAA
ATOM	2867	C	GLY	358					AAAA
ATOM	2868	0	GLY	358	9.626	40.757	71.663	1.00 24.03	
ATOM	2869	N	GLU	359	7.601	40.747	70.715	1.00 28.34	AAAA
				359	7.218	39.509	71.366	1.00 24.37	AAAA
MOTA	2870	CA	GLU				71.375	1.00 32.52	AAAA
MOTA	2871	CB	GLU	359	5.699	39.372			
MOTA	2872	CG	GLU	359	4.981	40:327	72.299	1.00 45.44	AAAA
ATOM	2873	CD	GLU	359	3.472	40.250	72.132	1.00 50.43	AAAA
					2.924	39.125	72.151	1.00 42.92	AAAA
MOTA	2874		GLU	359				1.00 40.72	AAAA
ATOM	2875	OE2	GLU	359	2.839	41.316	71.987		
MOTA	2876	C	GLU	359	7.804	38.323	70.628	1.00 27.35	AAAA
	2877	ō	GLU	359	8.138	38.415	69.449	1.00 22.94	AAAA
MOTA					7.944	37.208	71.325	1.00 19.68	AAAA
ATOM	2878	N	VAL	360					AAAA
ATOM	2879	CA	VAL	360	8.441	36.017	70.672	1.00 21.28	
ATOM	2880	СВ	VAL	360	9.300	35.188	71.621	1.00 26.71	AAAA
	2881		VAL	360	9.783	33.917	70.912	1.00 20.64	AAAA
ATOM					10.486	36.038	72.113	1.00 25.79	AAAA
MOTA	2882	CG2	VAL	360					AAAA
MOTA	2883	С	VAL	360	7.228	35.202	70.197	1.00 25.51	
MOTA	2884	0	VAL	360	6.442	34.700	71.011	1.00 19.75	AAAA
				361	7.065	35.094	ز 68.87	1.00 18.48	AAAA
MOTA	2885	N	ARG					1.00 22.01	AAAA
ATOM	2886	CA	ARG	361	5.947	34.337	68.3C`		
ATOM	2887	CB	ARG	361	5.988	34.389	66.772	1.00 19.31	AAAA
	2888	CG	ARG	361	5446	35.671	66.204	1.00 30.86	AAAA
MOTA					5.735	35.730	64.723	1.00 37.95	AAAA
ATOM	2889	CD.	ARG	361					AAAA.
MOTA	2890	NE	ARG	361	7.111	36.148	64.460	1.00 30.73	
MOTA	2891	CZ	ARG	361	7.616	36.275	63.242	1.00 22.89	AAAA
					6.851	36.006 [.]	62.186	1.00 19.02	AAAA
MOTA	2892		ARG	361				1.00 23.47	AAAA
ATOM	2893	NH2	ARG	3'61	8.861	36.704	63.081	1.00 23.47	
MOTA	2894	С	ARG	361	5.897	32.879	68.714	1.00 26.11	AAAA
				361	6.926	32.255	68.968	1.00 21.79	AAAA
ATOM	2895	0	ARG		4.681	32.338	68.763	1.00 24.89	AAAA
MOTA	2896	N	LYS	362					AAAA
ATOM	2897	CA	LYS	362	4.479	30.938	69.125	1.00 28.63	
ATOM	2898	CB	LYS	362	2.981	30.570	69.070	1.00 22.91	AAAA
					2.145	31.200	70.168	1.00 50.86	AAAA
ATOM	2899	CG	LYS	362				1.00 57.51	AAAA
ATOM	2900	CD	LYS	362	2.290		70.157	1.00 37.31	
ATOM	2901	CE	LYS	362	1.923	33.278	68.799	1.00 50.87	AAAA
				362	2.307	34.683	68.711	1.00 22.99	AAAA
ATOM .	2902	NZ	LYS			30.014	68.202	1.00 16.77	AAAA
MOTA	2903	С	LYS	362	5.269			1.00 10.00	
ATOM	2904	0	LYS	3.62	5.808	29.007	68.647	1.00 22.90	, AAAA
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Figure 16-45

	2025		OT 17	262		20 255	CC 013	1 00 05 04	
MOTA	2905	N	GLU		5.311	30.355	66.913	1.00 25.24	AAAA
MOTA	2906	ÇA	GLU	363	6.055	29.577	65.910	1.00 26.29	AAAA
ATOM	2907	СВ	GLU	363	6.207	30.342	64.608	1.00 33.50	AAAA
MOTA	2908	CG	GLU		4.999	30.639	63.824	1.00 48.73	AAAA
MOTA	2909	CD	GLU		5.368	31.494	62.638	1.00 42.01	AAAA
- MOTA	2910	OE1	L GLU	363	6.299	31.087	61.895	1.00 28.50	AAAA
MOTA	· 2911	OE2	GLU	363	4.738	32.558	62.461	1.00 44.91	AAAA
	2912				7.481	29.326	66.349	1.00 19.00	
ATOM		C	GLU						ÀAAA
MOTA	2913	0	GLU	363	8.011	28.218	66.226	1.00 18.66	AAAA
ATOM	2914	N	VAL	364	8.121	30.399	66.790	1.00 20.69	Aaaa
ATOM	2915	CA	VAL	364	9.501	30.303	67.219	1.00 23.13	AAAA
MOTA	2916	CB	VAL	.364	10.096	31.681	67.510	1.00 16.98	AAAA
MÓTA	2917	CG1	. VAL	364	11.515	31.513	68.010-	1.00 22.32	AAAA
ATOM	2918	CG2	. VAL	364	10.082	32.548	66.242	1.00 23.99	AAAA
ATOM	2919	C.	VAL	364	9.625	29.415	68.448	1.00 19.28	AAAA
ATOM	2920	0	VAL	364	10.507	28.548		1.00 20.17	- AAAA
MOTA	2921	N	LYS	365	8.735	29.600	69.417	1.00 21.11	AAAA
ATOM	2922	CA	LYS	365	8.780	28.768	70.612	1.00 18.15	AAAA
ATOM	2923	CB	LYS	365	7.711	29.210	71.626	1.00 25.22	AAAA
				365	7.921		72.167	1.00 32.99	
MOTA	2924	CG	LYS			30.611			AAAA
MOTA	2925	CD	LYS	365	6.901	30.949	73.253	1.00 36.09	AAAA
MOTA	2926	CE	LYS	365	7.121	32.357	73.790	1.00 28.99	AAAA
ATOM	2927	NZ	LYS	365	6.178	32.736	74.882	1.00 38.98	AAAA
			LYS	365	8.574	27.305	70.236	1.00 19.49	
MOTA	2928	C							AAAA
MOTA	2929	0	LYS	365	9.255	26.417	70.758	1.00 22.04	AAAA
MOTA	2930	N	ASP	3 6∙6	7.635	27.048	69.327	1.00 22.45	AAAA
ATOM	2931	CA	ASP	366	7.386	25.669	68.915	1.00 22.62	AAAA
MOTA	2932	CB	ASP	366	6.173	25.574	67.967	1.00 21.69	AAAA
MOTA	2933	CG	ASP	366	4.870	25.987	68.634	1.00 27.75	AAAA
MOTA	2934	OD1	ASP	366	4.763	25.890	69.881	1.00 31.01	AAAA
ATOM	2935	OD2	ASP	366	3.938	26.382	67.907	1.00 33.20	AAAA
ATOM	2936	C	ASP	366	8.606	25.034	68.237	1.00 24.53	AAAA
	2937				8.924	23.871	68.480		
MOTA		0	ASP	366				1.00 21.13	AAAA
ATOM	2938	N	THR	36 7	9.281	25.787	67.380	1.00 26.19	AAAA
ATOM	2939	CA	THR	367	10.462	25.252	66.694	1.00 21.68	AAAA
ATOM	2940	CB	THR	367	11.035	26.301	65.742	1.00 14.56	AAAA
	2941		THR	367	10.085	26.545	64.697	1.00 21.76	AAAA
ATOM									
ATOM	2942		THR	367	12.340	25.825	65.138	1.00 19.83	AAAA
ATOM	2943	С	THR	367	11.523	24.822	67.710	1.00 19.02	AAAA
ATOM	. 2944	0	THR	367	12.071	23.717	67.625	1.00 21.79	AAAA
ATOM	2945	N	LEU	368	11.802	25.684	68.683	1.00 18.42	AAAA
							69.700		
ATOM	2946	CA	LEU	368	12.797	25.348		1.00 21.02	AAAA
ATOM	2947	CB	LEU	368	13.148	26.569	70.560	1.00 17.34	AAAA
MOTA	2948	CG	LEU	368	14.206	27.518	69.959	1.00 17.45	AAAA
ATOM	2949	CD1	LEU	368	15.525	26.758	69.817	1.00 16.83	AAAA
	2950		LEU	368	13.756	28.041	68.593	1.00 19.49	AAAA
ATOM									
ATOM	2951	J	LEU	368	12.361	24.189	70.589	1.00 23.17	AAAA
ATOM	2952	U	LEU	368	13.203	23.420	71.052	1.00 24.81	AAAA
ATOM	2953	N	GLU	369	11.059	24.055	70.839	1.00 23.97	Aaaa
ATOM	2954	CA	GLU	369	10.597	22.929	71.653	1.00 19.36	AAAA
ATOM	2955	CB	GLU	369	9.127	23.113	72.063	1.00 21.81	AAAA
ATOM	2956	CG	GLU	369	8.913	24.225	73.100	1.00 40.15	AAAA
MOTA	2957	CD	GLU	369	7.450	24.416	73.487	1.00 49.38	AAAA
ATOM	2958		GLU	369	6.806	23.429	73.905	1.00 43.26	AAAA
MOTA	2959	OE2		369	6.948	25.558	73.382	1.00 57.31	AAAA
ATOM	2960	С	GLU	369	10.778	21.623	70.859	1.00 24.29	AAAA
ATOM	2961	9	GLU	369	11.172	20.605	71.420	1.00 25.96	AAAA
	2962	N	LYS	370	10.488	21.643	69.560	1.00 22.98	AAAA
MOTA							68.746	1.00 23.19	
ATOM	2963	CA	LYS	370	10.665	20.437			AAAA
MOTA	2964	CB .	LYS	370	10.051	20.596	67.347	1.00 26.83	AAAA
ATOM	2965	2G	LYS	370	8.537	20.461	67.287	1.00 36.68	AAAA
	2966	CD	LYS	370	8.056	20.431	65.832	1.00 39.85	AAAA
MOTA	_								
MOTA	2967	CE	LYS	370	6.567	20.105	65.740	1.00 56.23	AAAA
ATOM ·	2968	NZ	LYS	370	6.082	19.996	64.326	1.00 56.10	AAAA
MOTA	2969	С	LYS	370	12.148	20.123	68.602	1.00 31.63	Aaaa
ATOM	2970		LYS	370	12.549	18.958	68.587	1.00 36.88	AAAA
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ATOM	2971	N	ALA	371	12.961	21.170	68.491	1.00	26.25	AAAA
ATOM	2972		ALA	371	14.407		68.360		27.33	
										AAAA
MOTA	2973	CB	ALA	371	15.079	22.370	68.188	1.00	23.70	AAAA
ATOM	2974	С	ALA	371	14.989	20.308	69.581		26.74	AAAA
MOTA	2975	0	ALA	371	15.892	19.482	69.452	1.00	29.52	AAAA
ATOM	2976	N	ALA	372	14.484	20.652	70.764	1 00	24.83	AAAA
MOTA	2977	CA	ALA	372	14.959	20.055	72.012	1.00	34.24	AAAA
ATOM	2978	CB	ALA	372	14.305	20.750	73.214		37.17	
										AAAA
MOTA	2979	С	ALA	.372	14.663	18.564	72.061	1.00	45.62	AAAA
MOTA	2980	0	ALA	372	15.563	17.741	72.280		35.52	AAAA
MOTA	2981	N	ALA	373	13.394	18.216	71.869	1.00	44:50	AAAA
MOTA	2982	CA	ALA	373	13.004	16.813	71.892	1 00	49.88	AAAA
							_			
MOTA	2983	CB	ALA	373	11.506	16.681	71.628	1.00	49.32	AAAA
ATOM	2984	С	ALA	373	13.807	16.072	70.825	1.00	44.64	AAAA
MOTA	2985	0	ALA	373	14.669	15.250	71.201		58.19	AAAA
ATOM	2986	OXT	ALA	373	13.591	16.337	69.626	1.00	41.63	AAAA
HETATM				.1	36.368					
			TAW			43.907	49.242	1.00	13.03	SOLV
HETATM	2988	OH2	WAT	2	23.107	30.584	59.802	1.00	11.42	SOLV
HETATM			WAT	3	20.594	33.744				
							61.457		14.73	SOLV
HETATM	2990	OH2	WAT	4	31.359	16.551	51.590	1.00	19.84	SOLV
HETATM	7991	OHO	WAT	5	30.389	18.140	45.769		19.94	SOLV
HETATM	2992	OH2	$\mathbf{T}\mathbf{A}\mathbf{W}$	6	16.925	41.748	56.551	1.00	13.33	SOLV
HETATM	2993	OH2	$\mathbf{T}\mathbf{A}\mathbf{W}$	7	28.448	16.084	62.316	1 00	14.08	SOLV
HETATM	2994	OHZ	\mathbf{WAT}	8	40.375	38.476	55.678	1.00	19.10	SOLV
HETATM	2995	OH2	WAT	9	18.455	29.667	54.797	1.00	18.81	SOLV
					26.305		59.507			
HETATM			war	10		18.390	_		16.65	SOLV
HETATM	2997	OH2	WAT	11	50.145	32.063	58.142	1.00	16.53	SOLV
HETATM			WAT	12	45.935	30.996			25.08	
					•		40.672			SOLV .
HETATM	2999	OH2	TAW	13	26.358	43.110	74.179	1.00	22.91	SOLV
HETATM	3000	OHO	TAW	14	48.727	24.720	56.917		25.49	SOLV
HETATM	3001	OH2	TAW	15	30.244	18.663	50.165	1.00	25.78	SOLV
HETATM	3002	OH2	WAT	16	10.615	28.799	63.631	1 00	22.40	SOLV
HETATM	3003	OHZ	WAT	17	18.401	20.018	62.704	1.00	21.46	SOLV
HETATM	3004	OH2	WAT	18	22.195	47.791	60.896	1.00	26.19	SOLV
					3.278					
HETATM			WAT	19		32.141	65.350		20.38	SOLV
HETATM	3006	OH2	WAT	20	23.643	22.897	59.512	1.00	21.27	SOLV
HETATM	3007	OHO	WAT	21	50.287	23.101	48.818		19.73	SOLV
HETATM	3008	OHZ	WAT	22	44.725	34.256	46.541	1.00	18.74	SOLV
HETATM	3009	OHO	WAT	23	8.346	30.527	49.922	1 00	22.33	SOLV
HETATM	3010	OH2	WAT	24	39.855	33.795	67.390	1.00	20.43	SOLV
HETATM	3011	OH2	HAT	25	7.827	32.763	57.779	1.00	19.24	SOLV
		OH2			45.388	34.567				
HETATM				26			36.246		20.86	SOLV
HETATM	3013	OH2	WAT	27	47.636	32.244	33.388	1.00	20.41	SOLV
HETATM	3014	OH2	TAIL	28	32.514	35.684	41.278	1 00	24.76	SOLV
HETATM	3015	OH2	WAT	29	26.188	15.341	61.913	1.00	19.63	SOLV
HETATM	3016	OH2	WAT	30	14.957	43.169	56.333	1.00	23.80	SOLV
HETATM			WAT.	31	24.483	43.556	55.704	1.00	27.25	SOLV
HETATM	3018	OH2	WAT	32	41.141	16.376	48.456	1.00	25.99	SOLV
HETATM	7010	OH2		33	23.104	17.625	54.086		26.37	SOLV
HETATM	3020	OH2	WAT	34	51.301	28.602	57.694	1.00	32.78	SOLV
HETATM	3021	OH2	147 አጥ	35	51.376	29.469	53.156	1 00	24.27	SOLV
HETATM	3022	OH2	WAT	36	12.518	22.131	49.816	1.00	23.60	SOLV
HETATM	3023	OH2	WAT	37	6.521	27.442	50.861	1.00	25.87	SOLV
	2004									
HETATM	3024	OH2	MAT.	38	30.390	33.757	34.190		19.87	SOLV
HETATM	3025	OH2	WAT	39	8.328	29.586	62.062	1.00	32.01	SOLV
					30.180					
HETATM	3026	OH2		40		24.235	30.724		22.61	SOLV
HETATM	3027	OH2	WAT	41	44.521	30.663	38.395	1.00	27.52	SOLV
		OH2		42	30.981	18.043	41.186		23.45	SOLV
HETATM										
HETATM	3029	OH2	WAT	43	14.632	37.127	73.830	1.00	29.36	SOLV
HETATM		OH2		44	39.332	25.953	72.230		21.87	SOLV
HETATM	3031	OH2	WAT	45	7.597	37.592	51.896	1.00	39.62	SOLV
HETATM	3032	OH2	MAT	46	15.027	18.079	54.827	3 00	26.65	SOLV
UPIVIN	2024									
HETATM	3033	OH2	WAT	47	11.076	45.493	66.435		38.18	SOLV
HETATM	3034	OH2 1	MAT	48	42.124	18.055	37.233	1.00	28.62	SOLV
UPININ	2024						_			
HETATM	3035	OH2 1	WAT'	49	48.736	25.764	64.149	1.00		20TA
HETATM	3036	OH2		50	50.383	27.254	54.972	1.00	24.36	SOLV
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HETATM 3037	OH2 WAT	51	48.659	36.025	68.226	1.00 33.89	SOLV
HETATM 3038		52	36.998	27.228		1.00 21.03	SOLV
HETATM 3039	OH2 WAT	53	41.303	16.309	55.307	1.00 32.23	SOLV
HETATM 3040	OH2 WAT	54	33.242	39.524	49.454	1.00 29.77	SOLV
HETATM 3041		55	45.004	25.973		1.00 21.59	SOLV
HETATM 3042		56	19.039	25.829		1.00 33.48	SOLV
HETATM 3043	OH2 WAT	5 7	17.922	35.542	50.154	1.00 37.51	SOLV
HETATM 3044	OH2 WAT	58	10.409	26.864	73.166	1.00 26.54	SOLV
HETATM 3045		59	11.835	22.805			
						1.00 20.83	SOLV
HETATM 3046		60.	18.254	48.699		1.00 28.41	. SOLV
HETATM 3047	OH2 WAT	61	10.426	26.647	60.447	1.00 32.72	SOLV
HETATM 3048		62	21.304	55.086		1.00 28.84	SOLV
		63	32.532	51.211		1.00 32.48	
HETATM 3049							SOĻV
HETATM 3050	OH2 WAT	64	22.658	61.079	57.420	1.00 27.32	SOLV
HETATM 3051	OH2 WAT	65	16.734	24.334	74.721	1.00 27.44	SOLV
HETATM 3052	OH2 WAT	66	32.758	37.824	54.391	1.00 25.07	SOLV
HETATM 3053		67	11.142	25.859	49.706	1.00 29.66	
							SOLV
HETATM 3054		68	24.192	15.261	53.236	1.00 30.21	SOLV
HETATM 3055	OH2 WAT	69	19.816	17.916	66.357	1.00 30.50	SOLV
HETATM 3056	OH2 WAT	70 .	50.347	23.975	53.197	1.00 28.08	SOLV
HETATM 3057		71	50.258	30.918	51.113	1.00 20.19	
							SOLV
HETATM 3058	OH2 WAT	72	21.047	17.624	68.693	1.00 41.23	SOLV
HETATM 3059	OH2 WAT	73	26.782	33.756	49.995	1.00 25.80	SOLV
HETATM 3060	OH2 WAT	74	12.570	43.844	64.441	1.00 31.03	SOLV
HETATM 3061	OH2 WAT	75	35.555	41.287	50.852	1.00 24.03	SOLV
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нетатм 3062	OH2 WAT	76	27.764	18.231	61.827	1.00 18.28	SOLV
HETATM 3063	OH2 WAT	7 7	26.715	29.236	38.391	1.00 23.18	SOLV
HETATM 3064	OH2 WAT	78	21.461	23.245	48.872	1.00 23.80	SOLV
HETATM 3065	OH2 WAT	79	49.246	28.263	65.477	1.00 21.52	SOLV
			31.785				
HETATM 3066	OH2 WAT	80		13.301	69.606	1.00 31.11	SOLV
HETATM 3067	OH2 WAT	81	49.811	34.740	59.229	1.00 31.76	SOLV
HETATM 3068	OH2 WAT	82	45.670	33.188	42.470	1.00 23.13	SOLV
HETATM 3069	OH2 WAT	83	9.408	39.751	55.872	1.00 31.53	SOLV
			35.166				
HETATM 3070	OH2 WAT	84 .		35.878	29.899	1.00 37.32	SOLV
HETATM 3071	OH2 WAT	85	41.927	22.970	73.694	1.00 44.07	SOLV
HETATM 3072	OH2 WAT	86	22.125	34.577	49.199	1.00 44.65	SOLV
HETATM 3073	OH2 WAT	87	43.984	33.541	37.965	1.00 24.88	SOLV
HETATM 3074	OH2 WAT	88	11.997	17.962	56.312	1.00 34.85	SOLV
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HETATM 3075	OH2 WAT	89	42.194	14.737	59.766	1.00 25.91	SOLV
HETATM 3076	OH2 WAT	90	49.313	24.200	41.684	1.00 29.29	SOLV
HETATM 3077	OH2 WAT	91	48.504	33.595	61.519	1.00 30.32	SOLV
HETATM 3078	OH2 WAT	92	24.773	18.356	33.365	1.00 53.13	SOLV
		93	35.160		47.470		
HETATM 3079	OH2 WAT			35.656		1.00 41.41	SOLV
HETATM 3080	OH2 WAT	94	44.682	36.658	39.962	1.00 29.24	SOLV
HETATM 3081	OH2 WAT	95	. 9.576	41.033	52.549	1.00 51.83	· SOLV
HETATM 3082	OH2 WAT	^6	47.199	20.112	42.102	1.00 40.39	SOLV
HETATM 3083	OH2 WAT	7ر	49.254	26.331	59.641	1.00 37.03	SOLV
HETATM 3084	OH2 WAT	- 8	26.808	37.600	38.172	1.00 28.74	SOLV
HETATM 3085	OH2 WAT	99	40.749	14.572	64.635	1.00 33.42	SOLV
HETATM 3086	OH2 WAT	100	24.850	44.161	47.775	1.00 27.89	SOLV
HETATM 3087	OH2 WAT	101	34.326	42.063	46.714	1.00 42.22	SOLV
HETATM 3088	OH2 WAT	102	30.226	34.544	52.026	1.00 30.77	SOLV
HETATM 3089	OH2 WAT	103	47.824	39.054	78.097	1.00 52.16	SOLV
HETATM 3090	OH2 WAT	104	19.665	18.953	47.438	1.00 51.70	SOLV
HETATM 3091	OH2 WAT	105	46.857	36.525	46.232	1.00 23.65	SOLV
HETATM 3092	OH2 WAT	106	48.069	19.460	67.360	1.00 37.56	SOLV
HETATM 3093	OH2 WAT	107	15.553	56.850	61.838	1.00 46.95	SOLV
HETATM 3094	OH2 WAT	108	44.026	19.119	70.671	1.00 39.55	SOLV
HETATM 3095	OH2 WAT	109	8.139	42.064	65.674	1.00 42.61	SOLV
			50.624				
HETATM 3096	OH2 WAT	110		36.591	65.779	1.00 31.59	SOLV
HETATM 3097	OH2 WAT	111	51.398	26.073	61.043	1.00 49.09	SOLV
HETATM 3098	OH2 WAT	112	26.174	33.692	33.551	1.00 36.61	SOLV
HETATM 3099	OH2 WAT	113	23.545	20.203	53.001	1.00 24.34	SOLV
			9.083	42.965	57.697	1.00 33.65	SOLV
HETATM 3100	OH2 WAT	114					
HETATM 3101	OH2 WAT	115	8.442	39.898	64.594	1.00 31.21	SOLV
HETATM 3102	OH2 WAT	116	15.219	35.897	51.951	1.00 26.59	SOLV
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66/263 Figure 16-48

			i- 417	20 420	E0 455		
HETATM 310	3 OH2 WAT	117	15.417		50.473	1.00 34.46	SOLV
HETATM 310	4 OH2 WAT	118	40.757	26.310	29.206	1.00 29.12	SOLV
HETATM 310		119	27.717	18.542	46.553	1.00 28.17	SOLV
HETATM 310		. 120	18.612			1.00 38.56	SOLV
HETATM 310	7 OH2 WAT	121	43.198	31.377	72.139	1.00 26.31	SOLV
HETATM 310		122	44.188	35.704	33.802	1.00 29.81	SOLV
HETATM 310	9 OH2 WAT	123	50.736			1.00 32.40	SOLV
HETATM 311	O OH2 WAT	124	31.302	33.760	31.742	1.00 30.84	SOLV
				21.264			
HETATM 311	1 OH2 WAT	125	36.895			1.00 34.67	SOLV
HETATM 311	2 OH2 WAT	126	47.474	22.252	67.427	1.00 34.35	SOLV
HETATM 311		127	7.178	25.936	64.063	1.00 31.77	SOLV
HETATM 311	4 OH2 WAT	128	36.362	66.647	54.021	1.00 36.88	SOLV
HETATM 311	5 OH2 WAT	129	42.486	35.503	30.348	1.00 26.61	SOLV
HETATM 311		130	8.432	34.383	50.442	1.00 37.45	SOLV
HETATM 311	7 OH2 WAT	131	37.644			1.00 37.33	SOLV
HETATM 3118	B OH2 WAT	132	. 50.273	41.645	63.380	1.00 37.33	SOLV
		133	7.518	26.633	61.571	1.00 45.42	SOLV
HETATM 3119							
HETATM 3120	OH2 WAT	134	31.483	46.197		1.00 28.02	SOLV
HETATM 3123	L OH2 WAT	135	41.501	16.604	58.054	1.00 32.78	SOLV
		136	45.898	47.740	55.185	1.00 43.47	SOLV
HETATM 3122					33.103		
HETATM 3123	OH2 WAT	137	16.300	33.614	49.519	1.00 30.37	SOLV
HETATM 3124	OH2 WAT	138	51.148	36.946	55.148	1.00 46.34	SOLV
		139	21.525	53.761	50.892	1.00 38.27	SOLV
HETATM 3125					50.092		
HETATM 3126	OH2 WAT	140	21.603	54.580	68.690	1.00 33.10	SOLV
HETATM 3127	OH2 WAT	141	10.191	29.237	60.325	1.00 30.24	SOLV
HETATM 3128		142	16.951	18.120	66.901	1.00 40.85	SOLV
HETATM 3129	OH2 WAT	143	4.943	24.912	51.199	1.00 49.13	SOLV
HETATM 3130	OH2 WAT	144	10.711	25.291	58.177	1.00 30.72	SOLV
HETATM 3131		145	30.815	43.398	36.040	1.00 42.23	SOLV
HETATM 3132		146	21.763	24.512	46.695	1.00 28.31	SOLV
HETATM 3133	OH2 WAT	147	51.788	33.122	50.887	1.00 26.15	SOLV
HETATM 3134		148	24.531	44.741	72.420	1.00 27.99	SOLV
HETATM 3135		149	50.938		60.422	1.00 38.20	SOLV
HETATM 3136	OH2 WAT	150	24.860	47.932	61.067	1.00 18.89	SOLV
HETATM 3137		151	27.336	37.304	35.642	1.00 33.58	SOLV
			38.680	35.535	35.974	1.00 26.89	SOLV
HETATM 3138		152					
HETATM 3139	OH2 WAT	153	24.441	16.097	33.317	1.00 48.33	SOLV
HETATM 3140	OH2 WAT	154	20.343	18.124	73.416	1.00 36.28	SOLV
HETATM 3141		155	49.765	37.948	74.801	1.00 48.41	SOLV
HETATM 3142		156	34.329	31.169	47.547	1.00 25.33	sorv
HETATM 3143	OH2 WAT	157	43.028	24.554	72.536	1.00 41.54	SOLV
HETATM 3144		158	39.888	15.082	42.035	1.00 28.76	SOLV
HETATM 3145		159	41.886	20.780	73.179	1.00 51.03	SOLV
нетатм 3146	OH2 WAT	160	22.962	49.969	58.518	1.00 35.04	SOLV
HETATM 3147		161	14.696	15.261	68.016	1.00 55.47	SOLV
			14.915	18.181	64.866	1.00 42.00	
HETATM 3148		162					
HETATM 3149	OH2 WAT	163	30.608	49.029	52.612	1.00 47.32	. SOLV
HETATM 3150	OH2 WAT	164	52.566	30.906	57.612	1.00 36.71	SOLV
HETATM 3151	OH2 WAT	165	23.699	27.331	77.729	1.00 32.22	SOLV
HETATM 3152	OH2 WAT	166	36.971	59.046	63.272	1.00 43.05	SOLV
HETATM 3153	OH2 WAT	167	46.053	45.927	52.876	1.00 33.66	SOLV
HETATM 3154	OH2 WAT	168	42.780	49.151	58.106	.1.00 44.63	SOLV
HETATM 3155	OH2 WAT	169	15.100	44.506	72.183	1.00 45.43	SOLV
HETATM 3156	OH2 WAT	170	31.677	60.998	50.050	1.00 34.51	SOLV
HETATM 3157	OH2 WAT	171	25.336	45.674	45.578	1.00 55.85	SOLV
HETATM 3158	OH2 WAT	172	17.481	18.266	49.018	1.00 32.73	SOLV
HETATM 3159	OH2 WAT	173	26.112	18.147	31.404	1.00 49.94	SOLV
HETATM 3160	OH2 WAT	174	45.874	43.142	70.985	1.00 32.89	SOLV
HETATM 3161	OH2 WAT	175	34.517	17.884	33.278	1.00 42.20	SOLV
HETATM 3162	OH2 WAT	176	16.330	54.886	50.466	1.00 40.74	SOLV
HETATM 3163	OH2 WAT	177	31.400	51.087	74.689	1.00 38.56	SOLV
HEIMIN 3103							
HETATM 3164	OH2 WAT	178	50.971	27.079	67.130	1.00 44.49	SOLV
HETATM 3165	OH2 WAT	179	7.933	23.412	54.691	1.00 42.84	SOLV
	OH2 WAT	180	33.498	47.596	73.612	1.00 35.99	SOLV
HETATM 3166							SOLV
HETATM 3167	OH2 WAT	181	26.016	19.583	44.954	1.00 51.31	
HETATM 3168	OH2 WAT	182	40.139	17.026	74.920	1.00 43.64	SOLV
ADIAII 5100		- -	•		-		•

HETATM	3169	OH2	WAT	183	10.441	42.659	62.744	1.00 34.51	SOLV
HETATM	3170		TAW	184	2.095	34.482	65.810	1.00 36.49	SOLV
HETATM			WAT	185	45.749	18.286	51.615	1.00 28.19	SOLV
HETATM	3172		WAT	186	25.771	38.332	76.707	1.00 45.53	SOLV
HETATM	3173		WAT	187	7.228	40.382	57.542	1.00 48.91	SOLV
HETATM	3174		WAT	188	42.972	52.824	67.739	1.00 39.99	SOLV
HETATM	3175		WAT	189	20.137	13.189	73.277	1.00 44.91	SOLV
HETATM			TAW	190	48.945	19.193	47.581	1.00 52.88	.SOLV
HETATM	3177		TAW	191	14.549	34.547	47.665	1.00 49.15	SOLV
HETATM	3178		WAT	192	31.765	20.567	26.536	1.00 42.23	SOLV
HETATM	3179		WAT	193	9.784	39.303	74.222	1.00 32.10	SOLV
HETATM	3180		WAT	194	28.865	12.481	52.375	1.00 50.98	SOLV
HETATM	3181		WAT	195	24.030	12.804	70.409-	1.00 52.43	SOLV
HETATM	3182		WAT	196	47,209	39.53€	50.698	1.00 43.03	SOLV
HETATM	3183		WAT	197	35.618	18.114	27.306	1.00 41.11	SOLV
HETATM	3184		WAT	198	23.625	48.145	43.853	1.00 48.20	- SÓLV
HETATM	3185		WAT	199	37.090	59.044	54.185	1.00 34.99	SOLV
HETATM	3186		WAT	200	34.478	12.208	59.080	1.00 36.58	SOLV
HETATM	3187		WAT	201	22.142	29.583	76.228	1.00 33.95	· SOLV
HETATM	3188		WAT	202	13.608	42.619	53.973	1.00 40.44	SOLV
HETATM	3189		WAT	203	42.647	18.701	72.526	1.00 55.64	SOLV
HETATM	3190		WAT	204	37.005	35.993	77.480	1.00 34.82	SOLV
HETATM	3191		TAW	205	34.154	20.512	33.327	1.00 31.00	SOLV
HETATM	3192		WAT	206	37.264	57.546	47.642	1.00 49.58	SOLV
HETATM	3193		WAT	207	17.924	35.195	79.003	1.00 38.45	SOLV
HETATM	3194	OH2		208	51.172	31.581	62.378	1.00 35.37	SOLV
HETATM	3195	OH2		209	50.503	36.726	79.224	1.00 39.95	SOLV
HETATM	3196		WAT	210	18.382	13.162	63.852	1.00 52.08	SOLV
HETATM	3197		WAT	211	27.245	8.351	55.199	1.00 39.12	SOLV
HETATM	3198	OH2		212	18.354	13.545	59.540	1.00 30.15	SOLV
HETATM	3199	OH2		213	49.088	51.744	63.388	1.00 36.69	SOLV
HETATM	3200	OH2	WAT	214	23.251	33.160	50.871	1.00 42.11	SOLV
HETATM	3201	OH2	WAT	215	12.989	35.073	50.651	1.00 38.63	SOLV
HETATM	3202	OH2	WAT	216	24.414	44.460	43.239	1.00 37.93	SOLV
HETATM	3203	OH2	TAW	217	24.690	47.590	73.117	1.00 34.17	SOLV
HETATM		OH2	TAW	218	19.844	17.949	81.360	1.00 40.74	SOLV
HETATM		OHZ	WAT	219	40.169	27.215	74.247	1.00 37.83	SOLV
HETATM	3206	CH2	WAT	220	38.737	39.516	73.171	1.00 49.20	SOLV
HETATM	3207	OH2		221	50.628	21.408	46.879	1.00 45.57	SOLV
HETATM	3208	OH2	TAW	222	35.436	43.288	75.660	1.00 37.33	SOLV
HETATM	3209		TAW	223	34.390	16.963	55.285	1.00 35.10	SOLV
HETATM	3210	OH2		224	21.800	35.454	34.475	1.00 46.29	SOLV
HETATM	3211	OH2	TAW	225	15.751	40.989	46.787	1.00 62.75	SOLV
HETATM	3212		WAT	226	23.844	48.662	66.295	1.00 38.35	SOLV
HETATM	3213	OH2		227	47.225	20.562	55.117	1.00 49.99	SOLV
MELLYLLAM		OH2	VAT	228	23.426	19.272	50.565	1.00 30.07	SOLV

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					r	igure 17-1						
					Residue	∉ x	Y	Z	occ.	В	Segment	ID
. •	_					# A2 720	36.862	75.052		64.01	6	
MOTA	1	CB	ALA	A	2	43.739						
MOTA	2	С	АĻА	A	2	44.405.	38.106	72.971		60.02	6	
ATOM	3	0	ALA	Α	2	43.251	38.536	72.908	1.00	57.94	8	
			ALA		2	46.142	37.179	74.497	1.00	62.88	7	-
ATOM	4	N						73.923		63.02	6	
MOTA	5	CA	ALA	Α	2	44.776	36.966					
MOTA	6	N	LYS	Α	3	45.398	38.588	72.233		55.40	7	
ATOM	7	CA	LYS	Α	3	45.196	39.671	71.287	1.00	53.02	6	
	8	CB	LYS		3	46.443	39.830	70.421	1.00	53.11	6	
MOTA					-		40.093	71.217		57.36	6	
MOTA	9	CG	LYS			47.703						
ATOM	10	CD	LYS	Α	3	48.941	39.976	70.349		60.94	6	
ATOM .	11	CE	LYS	Α	3	48.909	40.957	69.196	1.00	63.48	6	
	12	NZ	LYS		3	50.075	40.765	68.294	1.00	66.87	7	
ATOM						43.986	39.401	70.399		50.49	6	
MOTA	1.3	С	LYS		3							
MOTA	14	0	LYS	А	3	43.691	38.255	70.063		52.50	8	
ATOM	15	N	VAL	Α	4	43.281	40.464	70.034	1.00	45.96	7	
			VAL		4	42.122	40.352	69.167	1.00	41.16	6	
ATOM	16	CA					41.272	69.638		41.53	6	
ATOM	17	CB	VAL		4	40.983						
MOTA	18	CG1	VAL	Α	4	39.734	41.028	68.797		40.07	6	
ATOM	19	CG2	VAL	Α	4	40.705	41.033	71.115	1.00	38.31	6	
			VAL		4	42.619	40.796	67.796	1.00	39.96	6`	
MOTA	20	C					41.914	67.645		39.15	8	
ATOM.	21	0	VAL	A	4	43.123						
ATOM	22	N	LYS	Α	5	42.486	39.916	66.807		36.24	7	
ATOM	23	CA	LYS	A	5	42.956	40.186	65.449	1.00	35.66	6	
			LYS		5	43.930	39.088	65.024	1.00	37.33	6	
MOTA	24	CB					38.978	65.860		38.24	6	
MOTA	25	CG	LYS		5	45.197						
MOTA	26	CD	LYS	A	5	46.113	40.179	65.659		35.41	6	
ATOM	27	CE	LYS	A	5	47.436	39.957	66.369	1.00	37.46	6	
	28	NZ	LYS		5	48.345	41.121	66.245	1.00	35.63	7	
MOTA						41.840	40.254	64.415		34.40		
ATOM	29	С	LYS		5							
MOTA	30	0	LYS	Α	5	40.788	39.641	64.588		33.92		
MOTA	31	N	LEU	A	6	42.082	40.983	63.329	1.00	32.52		
	32	CA	LEU	A	6	41.097	41.094	62.253	1.00	33.64	6	
MOTA			LEU		6	40.589	42.532	62.114	1.00	31.83	6	
MOTA	33	CB					42.823	61.248		32.93		
MOTA	34	CG	LEU		6	39.346					_	
MOTA	35	CD1	LEU	Α	6	39.356	44.295	60.899		28.95		
ATOM	36	CD2	LEU	Α	6	39.336	42.031	59.964	1.00	32.87	6	
	37	c	LEU		6	41.802	40.721	60.955	1.00	35.09	6	
MOTA					6	42.631	41.491	60.468		36.93		
MOTA	38	0	LEU					60.382		35.52		
MOTA	39	N	ILE	Α	7.	41.494	39.561				_	
MOTA	40	CA	ILE	Α	7	42.145	39.199	59.129	1.00	35.14		
ATOM	41	СВ	ILE		7	42.062	37.711	58.850	\cdot 1.00	33.68	6	
	42		ILE		7	42.731	37.409	57.517	1.00	32.87	6.	
ATOM						42.746	36.941	59.975		33.32		
ATOM	43		ILE		7						_	
ATOM	44	CD1	ILE	Α	7	42.744	35.451	59.755		35.09	_	
ATOM	45	С	ILE	A	7	41.487	39.935	57.971	1.00	37.13		
	46	Ō	ILE		7	40.258	39.933	57.855	1.00	35.21	. 8	
ATOM						42.304	40.563	57.124	1.00	37.25	. 7	
MOTA	47	N	GLY		8			55.994		38.69		
ATOM 1	48	CA	GLY	A	8	41.771	41.305					
ATOM	49	С	GLY	Α	8	42.809	41.939	55.079		39.73		
	50	0	GLY	_	8	44.015	41.827	55.321	1.00	39.21	. 8	
ATOM					9	42.335	42.622	54.033	1.00	39.41	. 7	
MOTA	51	N	THR					53.057		38.69		
ATOM	52	CA	THR	A	9	43.212	43.268					•
ATOM	53	CB	THR	Α	9	44.132	42.210	52.390		37.27		
	54		THR		9	44.754	42.771	51.230	1.00	36.82	. 8	
ATOM					9	43.332	40.972	52.001	1.00	38.59	6	
MOTA	55		THR				44.045	51.970		38.60	_	
ATOM	56	Ç	THR	A	9	42.447						
ATOM	57	0	THR		9	41.434	43.569	51.452		37.30		
	58	И	LEU		10	42.939	45.238	51.628	1.00	38.14		
ATOM					10	42.304	46.077	50.609	1.00	39.39	6	
MOTA	59	CA	LEU			43.026	47.418	50.456		38.98	_	
ATOM	60	CB	LEU		10							
ATOM	61	CG	LEU	A	10	42.836	48.506	51.510		39.68		
ATOM	62	CD1	LEU	Α	10	41.343	48.830	51.594	1.00	40.22		
			LEU		10	43.382	48.057	52.857	1.00	40.11	. 6	
ATOM	63						45.432	49.239		41.66	_	
ATOM	64	С	LEU		10	42.238					_	
ATOM	65	0	LEU	Α	10	41.462	45.863	48.381		42.08		
ATOM	66	N	ASP	Α	11	43.052	44.408	49.025	1.00	43.51	. 7	
A				-				-	•		•	
•		-										

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Figure 17-2

				_			42 671	42 721	40 222	4 00 47 0	-
ATOM	67	CA	ASF		11		43.071	43.731	47.737	1.00 47.2	
ATOM	68	CB	ASP	, A	11		44.250	42.765	47.694	1.00 51.0	3 6
MOTA	69	CG	ASP		11		45.579	43.479	47.858	1.00 54.1	.0 6
								44.282	46.975	1.00 55.9	
MOTA	70		ASP		11		45.944				
ATOM	71	OD2	ASP	, A	11		46.255	43.251	48.879	1.00 57.7	'98
ATOM	72	С	ASP		11		41.756	43.016	47.423	1.00 46.3	6 6
ATOM	73.	0	ASP	, A	11		41.472	42.702	46.266	1.00 43.4	
ATOM	74	N	TYR	A	12		40.954	42.767	48.456	1.00 46.8	0 7
	75	CA	TYR		12		39.654	42.116	48.284	1.00 45.9	2 6
ATOM											
ATOM	76	CB	TYR		12		38.953	41.942	49.638	1.00 41.3	
ATOM	77	CG	TYR	A	12		39.358	40.697	50.390	1.00 38.8	2 6
	78		TYR		12		39.531	40.720	51.775	1.00 37.9	5 6
ATOM											
ATOM	79	CE1	TYR	A	12		39.869	39.560	52.476	1.00 36.1	
ATOM	80	CD2	TYR	A	12		39.533	39.479	49.721	1.00 37.6	9 6
ATOM	81	CE2			12		39.868	38.316	50.415	1.00 35.8	3 6
ATOM	82	cz	TYR		12		40.032	38.365	51.787		
ATOM	83	OH	TYR	A	12		40.339	37.216	52.470	1.00 36.3	1 8
ATOM	84	С	TYR	2	12		38.786	42.966	47.378	1.00 46.5	6 6
									46.791		
ATOM	85	0	TYR		12		37.821	42.476		1.00 47.3	
ATOM	86	N	$GL\mathtt{Y}$	À	13		39.138	44.247	47.278	1.00 47.2	8 7
ATOM	87	CA	GLY	2	13		38.385	45.164	46.442	1.00 46.5	3 6
								44.934	44.968	1.00 45.6	
ATOM	88	С	GLY		13		38.650				
ATOM	89	0	GLY	Α	13		37.895	45.401	44.117	1.00 43.6	8 8
MOTA	90	N	LYS	A	14		39.725	44.210	44.672	1.00 46.5	2 7
							40.112	43.908	43.296	1.00 47.2	
MOTA	91	CA	LYS		14						
ATOM	92	CB	LYS	À	14		41.629	43.748	43.201	1.00 50.2	2 6
ATOM	93	CG	LYS	A	14		42.396	45.044	43.307	1.00 57.1	.2 6
	94	CD	LYS		14		42.038	46.004	42.161	1.00 63.6	
ATOM											
ATOM	95	CE	LYS	A	14		42.349	45.422	40.768	1.00 66.6	
- MOTA	96	NZ	LYS	Ą	14		41.529	44.220	40.387	1.00 67.7	0 7
ATOM	97	С	LYS		14		39.460	42.643	42.769	1.00 44.1	.8 6
							39.564		41.585	1.00 40.3	
MOTA	98	0	LYS		14			42.325			
MOTA	99	N	TYR	А	15		38.790	41.926	43.661	1.00 43.2	
ATOM	100	CA	TYR	A	15		38.145	40.665	43.317	1.00 43.1	.8 6
		•.					38.789	39.547	44.142	1.00 36.8	
MOTA	101	CB	TYR		15						•
MOTA	102	CG	TYR	Α	15		40.302	39.560	44.053	1.00 32.9	
MOTA	103	CD1	TYR	Α	15		41.084	39.107	45.108	1.00 30.9	0 6
		CE1	TYR		15		42.476	39.144	45.035	1.00 30.9	
ATOM	104										
ATOM	105	CD2	TYR	A	15		40.952	40.049	42.912	1.00 33.0	
MOTA	106	CE2	TYR	A	15		42.341	40.092	42.826	1.00 29.6	8 6
	107	CZ	TYR		15		43.098	39.639	43.890	1.00 30.9	9 6
ATOM											
ATOM	108	OH	TYR	À	15		44.471	39.673	43.809	1.00 28.0	
MOTA	109	С	TYR	À	15		36.661	40.778	43.621	1.00 45.5	6 6
ATOM	110	0	TYR	2	15		36.149	40.153	44.552	1.00 45.2	2 8
							35.981	41.599	42.830	1.00 48.8	
ATOM	111	N	ARG		16						
MOTA	112	CA	ARG	Α	16		34.553	41.819	42.999	1.70 53.2	
MOTA	113	CB	ARG	A	16		34.193	43.263	42.654	1' JO 57.1	1 5
	114	CG	ARG		16		34.852	44.330	43.490	.1 70 61.6	6 6
ATOM										1.00 67.0	
ATOM	115	CD	ARG	A	16		34.280	44.408	44.886		
ATOM	116	NE	ARG	A	16		34.798	45.590	45.569	1.00 73.5	
ATOM	117	CZ	ARG	2	16		34.612	46.837	45.141	1.00 75.0	3 6
						-				1.00 73.0	
ATOM		·NH1			16		33.917	47.065	44.033		
ATOM	119	NH2	ARG	Α	16		35.142	47.856	45.808	1.00 75.7	
ATOM	120	С	ARG	2	16		33.757	40.903	42.080	1.00 51.7	9 6
							34.192	40.593	40.970	1.00 50.8	
MOTA	121	0	ARG		16						
ATOM	122	N	TYR	A	17		32.596	40.463	42.552	1.00 50.1	
ATOM	123	CA	TYR		17		31.737	39.634	41.733	1.00 49.6	0 6
							30.534	39.119	42.528	1.00 45.8	
ATOM	124	CB	TYR		17						
ATOM	125	CG	TYR	Α	17		30.803	37.894	43.365	1.00 42.0	
ATOM	126	CD1	TYR	A	17		31.589	37.932	44.438	1.00 42.4	5 6
			TYR				31.960	36.780	45.193	1.00 43.3	
ATOM	127				17						
ATOM	128		TYR		17		30.185	36.680	43.062	1.00 39.8	
ATOM	129	CE2	TYR	A	17		30.443	35.526	43.803	1.00 40.5	
		CZ	TYR		17		31.333	35.578	44.869	1.00 41.9	8 6
ATOM	130								_	1.00 34.6	_
ATOM	131	OH	TYR	A	17		31.600	34.438	45.598		
ATOM	132	C	TYR	A	17		31.245	40.547	40.622	1.00 51.8	8 6
				-	٠.				-		

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. Figure 17-3

		_	msen a	. 17	31.332	41.772	40.726	1.00 47.86	8
ATOM	133	0	TYR A				39.534	1.00 54.38	7
ATOM	134	N	PRO A	. 18	30.730	39.964			
ATOM	135	CD	PRO A	. 18	30.548	38.545	39.190	1.00 54.21	6
	136	CA.	PRO A		30.243	40.809	38.449	1.00 56.43	6
MOTA	_				29.601	39.792	37.496	1.00 56.84	6
ATOM	137	CB	PRO A	_		38.613	38.426	1.00 56.46	6
ATOM -	138	CG	PRO A		29.260				6
MOTA	139	С	PRO A	. 18	29.273	41.891	38.932	1.00 58.74	
ATOM	140	0	PRO A	. 18	28.791	41.861	40.066	1.00 55.72	8
			LYS A		29.017	42.851	38.052	1.00 62.10	7
MOŢA	141	N				43.973	38.314	1.00 64.85	6
ATOM	142	CA	LYS A		28.127				
MOTA	143	CB	LYS A	. 19	27.972	44.781	37.022	1.00 69.74	6
MOTA	144	CG	LYS A	. 19	28.008	43.925	35.740	1.00 74.99	6
	145	CD	LYS A		- 26.895	42.881	35.568-	1.00 78.18	6
ATOM					26.981	42.010	34.420	1.00 80.24	- 6
MOTA	146	CE	LYS A			41.010		1.00 81.13	7
MOTA	147	NZ	LYS A	. 19	25.867		34.361		
ATOM	148	С	LYS. A	. 19	26.750	43.619	38.869	1.00 64.77	6.
ATOM	149	0	LYS A	. 19	26.414	43.961	40.001	1.00 66.50	.8
			ASN A		25.957	42.933	38.062	1.00 63.75	7
MOTA	150	N			24.612	42.556	38.439	1.00 62.96	6
ATOM	151	CA	ASN A						6
ATOM	152	CB	ASN A	. 20	23.870	42.031	37.208	1.00 67.42	
ATOM	153	CG	ASN A	. 20	22.392	41.833	37.459	1.00 72.29	6
	154		ASN A		21.666	42.785	37.772	1.00 75.25	8
MOTA					21.931	40.594	37.322	1.00 74.38	7
ATOM	155		ASN A			41.512	39.547	1.00 61.30	6
ATOM	156	С	ASN A	20	24.602				
ATOM	157	0	ASN A	. 20	23.629	40.773	39.698	1.00 61.49	8
ATOM	158	N	HIS A	21	25.681	41.444	40.321	1.00 57.30	7
	159	CA	HIS A		25.755	40.480	41.418	1.00 54.68	6
ATOM					27.071	39.700	41.373	1.00 52.63	6
MOTA	160	CB	HIS A			38.449	42.195	1.00 49.39	6
ATOM	161	CG	HIS A	. 21	27.058				6
MOTA	162	CD2	HIS A	. 21	27.336	38.236	43.503	1.00 49.39	
ATOM	163	ND1	HIS A	21	26.664	37.229	41.686	1.00 48.27	7
	164		HIS A		26.704	36.320	42.643	1.00 48.16	6
MOTA					27.108	36.905	43.757	1.00 47.33	7
ATOM	165		HIS A				42.760	1.00 52.89	6
MOTA	166	С	HIS A	. 21	25.664	41.215			
MOTA	167	0	HIS A	21	26.295	42.256	42.947	1.00 52.52	8
ATOM	168	N	PRO A	22	24.880	40.679	43.713	1.00 50.81	7
	169	CD	PRO A		24.076	39.444	43.661	1.00 48.50	6
MOTA					24.734	41.310	45.029	1.00 48.02	6
MOTA	170	CA	PRO A		•		45.783	1.00 47.45	6
ATOM	171	CB	PRO A	. 22	23.860	40.308			
MOTA	172	CG	PRO A	. 22	22.990	39.754	44.667	1.00 47.76	6
MOTA	173	С	PRO A	. 22	26.074	41.558	45.727	1.00 46.48	6
	174	ŏ	PRO A		26.164	42.405	46.615	1.00 45.69	8
MOTA					27.107	40.816	45.318	1.00 44.97	7
ATOM	175	N	LEU A				45.906	1.00 41.31	6
MOTA	176	CA	LEU A		28.441	40.949			6
ATOM	177	CB	LEU A	. 23	29.076	39.569	46.131	1.00 39.22	
ATOM	178	CG	LEU A	23	28.264	38.561	46.953	1.00 37.71	6
	179		LEU A		29.075	37.288	47.157	1.00 35.07	6
MOTA					27.896	39.165	48.292	1.00 36.90	6
MOTA	180		LEU A				45.003	1.00 40.14	6
ATOM	181	Ç.	LEU A	. 23	29.334	41.789			8
ATOM	182	0	LEU A	23	30.556	41.614	44.951	1.00 39.00	
	183	N	LYS A	24	28.706	42.705	44.284	1.00 39.67	7
MOTA				- 4	29.430	43.590	43.398	.1.00 42.88	6
ATOM	184	CA	LYS A		28.480	44.120	42,323	1.00 40.24	6
ATOM	185	CB	LYS A					1.00 44.08	6
ATOM	186	CG	LYS A	24	28.949	45.362	41.610		6
ATOM	187	CD	LYS A	24	28.247	46.600	42.166	1.00 44.47	
	188	CE	LYS A		26.732	46.492	41.968	1.00 43.23	6
ATOM					25.989	47.717	42.362	1.00 39.79	7
ATOM	189	NZ	LYS A			44.723	44.217	1.00 43.70	6
ATOM	190	С	LYS A		30.031			1.00 47.22	8
ATOM	191	0	LYS A	24	31.027	45.332	43.817		7
ATOM	192	N	ILE A		29.431	44.976	45.378	1.00 42.27	7
		.,			29.870	46.035	46.289	1.00 39.86	6
		C 3	TITE N						
ATOM	193	CA	ILE A	_	20 762	46 407	47 306	1.00 37.72	6
		CA CB	ILE A	25	28.763	46.407	47.306	1.00 37.72	6 6
atom Atom	193 194			25	27.539	46.953	46.580	1.00 39.67	6
ATOM ATOM ATOM	193 194 195	CB CG2	ILE A	25		46.953 45.168	46.580 48.145	1.00 39.67 1.00 35.25	6 6
atom atom atom atom	193 194 195 196	CB CG2 CG1	ILE A	25 25 25	27.539 28.410	46.953	46.580 48.145 49.151	1.00 39.67 1.00 35.25 1.00 31.74	6 6
ATOM ATOM ATOM	193 194 195	CB CG2 CG1	ILE A	25 25 25 25	27.539	46.953 45.168	46.580 48.145	1.00 39.67 1.00 35.25	6 6

		_			24 410	44 449	45 100	4 60 30 00	_
MOTA	199	0	ILE A	25	31.419	44.441	47.198	1.00 38.90	8
ATOM	200	N	PRO A	26	31.762	46.616	47.709	1.00 40.18	7
ATOM	201	CD	PRO A	26	31.523	48.051	47.533	1.00 40.58	6
ATOM	202	CA	PRO A	26	32.939	46.437	48.558	1.00 38.31	6
	203		PRO A		33.478	47.860	48.688	1.00 37.14	6
ATOM		CB		26					
MOTA	204	CG	PRO A	26	32.940	48.537	47.458	1.00 38.77	6
	205	С	PRO A	26	32.433	45.903	49.891	1.00 37.32	6
ATOM									
MOTA	206	0	PRO A	26	31.416	46.372	50.412	1.00 32.70	8
ATOM	207	N	ARG A	27	33.134	44.930	50.452	1.00 36.54	7
			• •			44.359	51.711	1.00 37.39	6
ATOM	208	CA	ARG A	27	32.685				
ATOM	209	CB	ARG A	27	32.116	42.952	51.455	1.00 35.29	6
	210	CG	ARG A	27	31.047	42.956	50.355	1.00 32.69	б
MOTA								•	
MOTA	211	CD	ARG A	27	30.507	41.573	49.956	1.00 33.87	6
ATOM	212	NE	ARG A	27	29.757	40.909	51.021	1.00 36.16	7
. –		cz	ARG A		30,293	40.132	51.959	1.00 37.11	6
ATOM	213		-	27	-				
ATOM	214	NH1	ARG A	27	31.604	39.903	51.976	1.00 34.42	7
ATOM	215	NH2	ARG A	27	29.516	39.597	52.896	1.00 33.67	7
						44.329	52.732	1.00 36.35	6
ATOM	216	С	ARG A	27	33.813				
ATOM	217	0	ARG A	27	33.881	45.188	53.610	1.00 35.77	8
ATOM	218	N	VAL A	28	34.703	43.351	52.607	1.00 34.93	7
					35.810	43.230	53.537	1.00 34.00	6
ATOM	219	CA	VAL A	28					
ATOM	220	CB	VAL A	28	36.633	41.954	53.252	1.00 36.21	6
MOTA	221	CG1	VAL A	28	37.574	41.652	54.424	1.00 33.59	6
						40.790	52.992	1.00 37.05	6
MOTA	222	CG2		28	35.696				
ATOM	223	С	VAL A	28	36.712	44.454	53.423	1.00 31.91	6
ATOM	224	0	VAL A	28	37.216	44.959	54.427	1.00 31.45	8
					36.908	44.936	52.199	1.00 33.12	
MOTA	225	N	SER A	29					
ATOM	226	CA	SER A	29	37.751	46.111	51.9 67	1.00 32.03	6
ATOM	227	CB '	SER A	· 29	38.205	46.181	50.499	1.00 31.77	6
					37.113	46.223	49.600	1.00 30.80	8
MOTA	228	OG	SER A	29					
MOTA	229	С	SER A	29	37.003	47.380	52.353	1.00 30.16	6
ATOM	230	0	SER A	29	37.604	48.404	52.650	1.00 28.70	8
	231	N	LEU A	30	35.682	47.310	52.352	1.00 32.43	7
ATOM									
MOTA	232	CA	LEU A	30	34.900	48:465	52.745	1.00 34.56	6
ATOM .	233	CB	LEU A	30	33.463	48.358	52.221	1.00 36.44	6
	234	CG	LEU A	30	32.508	49.513	52.560	1.00 36.79	6
MOTA									
ATOM	235	CDI	LEU A	30	32.070	49.446	54.012	1.00 36.73	6
ATOM	236	CD2	LEU A	30	33.202	50.840	52.256	1.00 37.84	6
					34.902	48.527	54.262	1.00 34.89	6
MOTA	237	С	LEU A	30					
ATOM	238	0	LEU A	30	35.033	49.601	54.841	1.00 37.58	8
ATOM	239	N	LEU A	31	34.761	47.366	54.897	1.00 34.07	7
	240	CA	LEU A	31	34.743	47.276	56.350	1.00 34.85	6
ATOM					-				
ATOM	241	CB	LEU A	31	34.768	45.808	56.791	1.00 36.37	6
MOTA	242	CG	LEU A	31	34.459	45.471	58.261	1.00 36.04	6
	243			31	34.841	44.027	58.532	1.00 35.13	6
ATOM			LEU A						
ATOM	244	CD2	LEU A	31	35.228	46.357	59.194	1.00 35.86	. 6
MOTA	245	С	LEU A	31	35.976	47.994	56.894	1.00 36.43	. 6
	246	0	LEU A	31	35.855	49.035	57.544	1.00 35.87	8
ATOM									7
ATOM	247	N	LEU A	32	37.157	47.426	56.635	1.00 37.76	
ATCM	248	CA	LEU A	32	38.420	48.015	57.087	1.00 36.82	ő
	249	CB	LEU A	32	39.611	47.318	56.418	1.00 36.37	6
ATOM					<u> </u>				6
ATOM	250	CG	LEU A	32	40.030	45.888	56.774	1.00 39.11	
ATOM	251	CD1	LEU A	32	41.117	45.420	55.815	1.00 35.16	6
	252		LEU A	32	40.538	45.830	58.214	1.00 37.73	6
ATOM									6
ATOM	253	С	LEU A	32	38.500	49.513	56.780	1.00 34.84	
ATOM	254	0	LEU A	32	38.846	50.326	57.644	1.00 36.58	8
	255		ARG A	33	38.184	49.877	55.545	1.00 31.37	7
ATOM		N							6
ATOM	256	CA	ARG A	33	38.247	51.270	55.150	1.00 32.53	
MOTA	257 °	CB	ARG A	33	. 37.927	51.398	53.662	1.00 31.52	6
	258		ARG A	33	38.481	52.652	53.042	1.00 35.88	6
ATOM		CG							6
ATOM	259	CD	ARG A	33	38.107	52.752	51.581	1.00 43.44	-
ATOM	260	NE	ARG A	33	38.521	51.583	50.811	1.00 48.37	7
	261	CZ	ARG A	33	38.348	51.469	49.497	1.00 52.27	6
ATOM									7
ATOM	262		ARG A	33	37.771	52.459	48.823	1.00 51.75	
ATCM	263	NH2	ARG A	33	38.739	50.369	48.858	1.00 51.08	7
	264	C	ARG A	33	37.274	52.102	55.989	1.00 32.32	6
ATCM	204	_			↓ , 2			~ ~~~~~	

	265	_	א ספא	33		37.471	53.299	56.196	1.00 29.23	8
MOTA	265	0	ARG A					56.484	1.00 32.58	7
ATOM	266	N	PHE A			36.231	51.445			
ATOM	267	CA	PHE A	34		35.216	52.096	57.304	1.00 32.69	6
	268	CB	PHE A	34		33.952	51.232	57.359	1.00 31.22	6
ATOM			PHE A			32.838	51.825	58.183	1.00 28.74	6
MOTA	269	CG					52.888	57.700	1.00 22.76	6
ATOM	270	CD1	PHE A			32.085				
MOTA	271	CD2	PHE A	34		32.551	51.322	59.456	1.00 28.09	6
	272		PHE A	34		31.061	53.441	58.472	1.00 23.70	6
ATOM						31.524	51.873	60.235	1.00 24.59	6
MOTA	273	CE2	PHE A				52.929	59.741	1.00 21.39	6 -
ATOM	274	CZ	PHE A			30.781				
ATOM	275	С	PHE A	34		35.734	52.319	58.719	1.00 33.45	6
	276	0	PHE A	34		35.635	53.425	59.258	1.00 35.49	8
ATOM			LYS A			36.276	51.264	59.323	1.00 34.52	7
ATOM	277	N				36.805	51.360	60.678	1.00 36.51	6
MOTA	278	CA	LYS A	_					1.00 36.47	
MOTA	279	CB	LYS A	. 35		37.118	49.977	61.235		6
MOTA	280	CG	LYS A	35		35.912	49.074	61.343	1.00 40.81	6
	281	CD	LYS A			36.246	47.801	62.090	1.00 44.10	6
ATOM				-		37.347	47.029	61.402	1.00 47.46	6
MOTA	282	CE	LYS A					61.276	1.00 53.53	7
ATOM	283	NZ	LYS A	35		38.601	47.823			
ATOM	284	С	LYS A	35		38.054	52.222	60.735	1.00 36.61	6
	285	0	LYS A	35		38.352	52.824	61.766	1.00 36.78	8
MOTA			ASP A			38.794	52.267	59.635	1.00 36.27	7
MOTA	286	N					53.090	59.592	1.00 39.71	6
MOTA	287	CA	ASP A			39.980				6
MOTA	288	CB	ASP A	36		40.679	52.937	58.239	1.00 44.78	
MOTA	289	CG	ASP A	36		41.863	53.892	58.075	1.00 47.10	6
	290		ASP A	36		42.803	53.852	58.906	1.00 44.02	8
MOTA			ASP A			41.843	54.682	57.106	1.00 48.43	8
MOTA	291						54.530	59.789	1.00 39.99	6
MOTA	292	С	ASP A		٠.	39.508			1.00 40.76	8
ATOM	293	0	ASP A	36		40.023	55.258	60.636		
ATOM	294	N	ALA A	37		38.506	54.919	59.007	1.00 38.59	7
	295	CA	ALA A			37.939	56.258	59.066	1.00 37.14	6
MOTA						36.857	56.402	58.000	1.00 35.85	6
ATOM	296	CB	ALA A					60.446	1.00 38.34	6
ATOM	297	С	ALA A			37.354	56.549			
MOTA	298	0	ALA A	37		37.391	57.687	60.928	1.00 37.32	8
MOTA	299	N	MET A	38		36.809	55.518	61.079	1.00 36.19	7
	300	CA	MET A			36.213	55.674	62.397	1.00 36.80	6
ATOM						35.141	54.598	62.606	1.00 37.38	5
ATOM	301	CB	MET A					61.673	1.00 37.60	6
MOTA	302	CG	MET A			33.938	54.717			16
MOTA	303	SD	MET A	38		32.887	56.165	61.999	1.00 33.61	
ATOM .	304	CE	MET A	38		32.398	55.824	63.680	1.00 35.60	6
	305	c	MET A			37.262	55.582	63.502	1.00 35.84	6
MOTA			MET A			36.937	55.688	64.692	1.00 34.89	8
MOTA	306	0					55.400	63.100	1.00 33.83	7
ATOM	307	N	ASN A			38.518				6
ATOM	308·	CA	ASN A	39		39.626	55.264	64.044		
MOTA	309	CB	ASN A	39		39.897	56.582	64.775	1.00 32.48	6
	310	CG	ASN A			40.213	57.717	63.825	1.00 32.34	6
ATOM					•	41.128	57.621	63.009	1.00 31.85	8
MOTA	311		ASN A					63.924	1.00 30.92	7
ATOM	312		ASN A			39.455	58.800			6
ATOM	313	С	ASN A	39		39.253	54.183	65.045	1.00 36.87	
ATOM	314	0	ASN A	39		39.403	54.357	66.260	1.00 36.60 [.]	8
	315	N	LEU A			38.752	53.067	64.518	1.00 37.48	7
MOTA						38.341	51.933	65.336	1.00 39.66	6
ATOM	316	CA	LEU A						1.00 41.35	6
ATOM	317	CB	LEU A	40		36.863	51.622	65.086	1.00 41.33	
ATOM	. 318	CG	LEU A	40		35.858	52.712	65.476	1.00 42.69	6
	319		LEU A			34.448	52.261	65.111	1.00 45.05	6
ATOM						35.951	52.989	66.966	1.00 39.44	6
MOTA	320		LEU A	_			50.687	65.058	1.00 39.79	6
ATOM	321	С	LEU A			39.184			1.00 36.88	. 8
ATOM	322	0	LEU A	40		38.804	49.575	65.434		
	323	N	ILE A			40.337	50.889	64.420	1.00 40.50	7
MOTA		CA	ILE A			41.237	49.790	64.068	1.00 41.39	6
MOTA	324						49.141	62.724	1.00 39.24	6
ATOM	325	СВ	ILE A	_		40.780			1.00 36.97	6
ATOM	326	CG2	ILE A	41		41.017	50.103	61.564		
MOTA	327	CG1	ILE A	41		41.513	47.824	62.482	1.00 36.76	6
	328		ILE A			41.085	46.715	63.403	1.00 35.59	6
MOTA						42.684	50.295	63.913	1.00 44.37	6
ATOM	329	C	ILE A			_		63.277	1.00 46.01	8
ATOM	330	0	ILE A	41		42.927	51,328	55.277	2.00	-

MOTA	331		ASP .			43.646	49.582	64.497	1.00 45.19	7
MOTA	332	CA	ASP .	A 42		45.049	49.982	64.372	1.00 45.62	6
MOTA	333	CB	ASP :	A 42		45.716	50.090	65.742	1.00 45.17	6
ATOM	334	L CG	ASP 2			44.966	51.005	66.682	1.00 44.43	
ATOM	335		1 ASP							6
						44.731	52.177	66.322	1.00 39.84	8
ATOM	336					44.612	50.546	67.787	1.00 48.50	8
ATOM	337	, c	ASP	A 42		45.750	48.915	63.551	1.00 48.47	6
MOTA	338	0	ASP 2	A 42		45.316	47.757	63.547	1.00 49.85	8
ATOM	339	N	GLU Z			46.830	49.288	62.864	1.00 49.24	
ATOM	340		GLU A							7
						47.553	48.325	62.028	1.00 50.79	6
ATOM	341		GLU A			. 48.820	48.956	61.431	1.00 49.90	6
ATOM	342		GLU A	43		48.544	50.029	60.378	1.00 57.20	6
ATOM	343	CD	GLU A	A 43		49.808	50.537	59.690	1.00 59.56	6
ATOM	344	OE	L GLU A	4 43		50.517	49.721	59.061	1.00 65.05	8
ATOM	345	OE2	GLU A	4 43		50.095	51.750	59.772	1.00 57.82	
ATOM	346		GLU 7			47.918	47.020			8
	347	-						62.733	1.00 49.73	6
ATOM			GLU A			47.813	45.943	62.149	1.00 49.18	8
ATOM	348		LYS A			48.324	47.118	63.992	1.00 49.40	7
MOTA	349		LYS A	44		48.730	45.949	64.762	1.00 49.09	6
ATOM	350	CB	LYS A	44		49.317	46.418	66.093	1.00 52.46	6
ATOM	351	CG	LYS A	44		50.448	47.421	65.899	1.00 55.75	6
ATOM	352	CD	LYS A			51.167	47.749	67.201	1.00 58.74	
ATOM	353	CE	LYS A			52.327	48.704	66.949		6
ATOM	354	NZ	LYS A						1.00 58.48	6
						53.122	48.968	68.176	1.00 58.95	7
MOTA	355	C	LYS A			47.638	44.897	64.994	1.00 47.63	6
ATOM	356	0	LYS A			47.932	43.738	65.290	1.00 45.13	8
MOTA	357	N	GLU A			46.379	45.298	64.854	1.00 45.25	7
MOTA	358	CA	GLU A	. 45		.45.268	44.374	65.046	1.00 43.09	б
ATOM	359	CB	GLU A	45		44.024	45.143	65.514	1.00 41.19	6
ATOM-	360	ÇG	GLU A	45		44.192	45.859	66.844	1.00 36.83	6
ATOM	361	CD	GLU A			43.003	46.741	67.204	1.00 38.92	
ATOM	362		GLU A			42.707				6
	363						47.701	66.447	1.00 37.30	8
MOTA			GLU A			42.368	46.479	68.253	1.00 36.33	8
ATOM	364	Ç	GLU A			44.969	43.660	63.726	1.00 43.04	6
MOTA	. 365.	0	.GLU A			44.480			1.00 45.03	8
ATOM	366	N	LEU A	46		45.282	44.341	62.632	1.00 40.29	7.
ATOM	367	CA	LEU A	46		45.042	43.823	61.299	1.00 37.16	6
ATOM	368	CB	LEU A	46		44.910	44.990	60.331	1.00 37.86	6
ATOM	369	CG	LEU A			44.822	44.658	58.845	1.00 39.22	
MOTA	370		LEU A			43.655				6
	371		LEU A				43.726	58.563	1.00 40.68	6
ATOM						44.673	45.964	58.080	1.00 41.62	6
ATOM	372	С	LEU A			46.090	42.860	60.774	1.00 36.54	6
ATOM	373	0	LEU A			· 47.275	43.192	60.698	1.00 39.86	8
MOTA	374	N	ILE A	47		45.646	41.662	60.406	1.00 33.49	7
MOTA	375	CA	ILE A	47		46.540	40.657	59.844	1.00 30.51	6
ATOM	376	CB	ILE A	47		46.333	39.253	60.491	1.00 34.31	6
ATOM	377	CG2		47		47.346	38.262	59.930	1.00 32.16	6
ATOM	378	CG1	ILE A	47		46.504	39.328	62.010	1.00 32.65	
ATOM	379		ILE A							6
					· ·	.47.858	39.846	62.448	1.00 38.97	6
MOTA	380	C	ILE A	47		46.196	40.570	58.362	1.00 28.36	6
MOTA	381	0	ILE A	47		45.037	40.342	58.003	1.00 26.11	8
ATOM	382	N	LYS A	48		47.194	40.772	57.504	1.00 27.77	7
ATOM	383	CA	LYS A	48		46.985	40.713	56.056	1.00 25.80	6
MOTA	384	CB	LYS A	48		48.258	41.087		1.00 23.91	6
MOTA	385	CG	LYS A	48		48.056	41.273	53.811	1.00 24.90	6
ATOM	386	CD	LYS A	48		49.389	41.352	53.091	1.00 26.39	
	387									6
ATOM		CE	LYS A	48		49.233	41.864	51.679	1.00 27.71	6
ATOM	388	NZ	LYS A	48		48.774	43.275	51.696	1.00 32.59	7
MOTA	389	C	LYS A	48		46.595	39.299	55.654	1.00 26.32	6
ATCM	390	0	LYS A	48		47.072	38.325	56.235	1.00 27.85	8
ATOM	391	N	SER A	49		45.735	39.183	54.653	1.00 24.73	7
MOTE	392	CA	SER A	49		45.299	37.876	54.205	1.00 27.36	6
MOTA	393	CB	SER A	49		43.952	37.979	53.479	1.00 25.04	6
	394	OG	SER A	49	٠.	42.911	38.329	54.373	1.00 26.94	
MOTA				49						8
MOTA	395	C	SER A			46.322	37.211	53.293	1.00 28.97	6
ÄTOM	396	0	SER A	49		47.095	37.885	52.612	1.00 31.89	8
		•		•	•				•	

ATOM	397 398	N CA	ARG A ARG A	50		46.315	35.879 35.087	53.296	1.00 29.71	7
ATOM ATOM	399	CB	ARG A	50 50		47.211 48.249	34.351	52.463 53.318	1.00 25.78 1.00 26.20	6 6
ATOM	400	CG	ARG A	50		47.687	33.204	54.172	1.00 22.71	6
ATOM	401	CD	ARG A	50		48.818	32.468	54.890	1.00 22.95	6
ATOM_	402	NE	ARG A	50		48.359	31.385	55.762	1.00 19.20	7
MOTA	403	CZ	ARG A	50		47.708	30.306	55.345	1.00 16.85	6
ATOM	404		L ARG A	50		47.430	30.151	54.055	1.00 17.77	7
ATOM	405	NH2		50		47.334	29.385	56.223	1.00 14.56	7
ATOM	406	C	_ARG A	50		46.370	34.051	51.723	1.00 23.30	6
MOTA	407 408	N O	ARG A PRO A	50 51		45.319 46.823	33.635 33.628	52.206 50.534	1.00 16.92	8 7
ATOM ATOM	409	CD	PRO A	51		48.021	34.038	49.789	1.00 21.06 1.00 20.50	6
MOTA	410	CA	PRO A	51		46.086	32.633	49.761	1.00 20.50	6
ATOM	411	CB.	PRO A	51		46.862	32.592	48.451	1.00 21.57	6
MOTA	412	CG	PRO A	51		47.503	33.984	48.392	1.00 20.57	6.
MOTA	413	C	PRO A	51		46.153	31.300	50.498	1.00 26.71	6
ATOM	414	0	PRO A	51	,	47.071	31.066	51.293	1.00 31.32	8
MOTA	415	N	ALA A	52		45.176	30.435	50.250	1.00 26.02	7
MOTA	416 417	CA CB	ALA A ALA A	52 52		45.151 43.720	29.121 28.585	50.876 50.933	1.00 25.76	6
ATOM ATOM	418	C	ALA A	52		46.013	28.227	50.000	1.00 21.42 1.00 26.31	6 6
ATOM	419	ō	ALA A	52		45.878	28.239	48.780	1.00 20.31	. 8
ATOM	420	N	THR A	53		46.909	27.464	50.608	1.00 26.80	. 7
MOTA	421	CA	THR A	53		47.759	26.578	49.831	1.00 27.52	6
ATOM	422	CB	THR A	53		48.845	25.975	50.71 7	1.00 26.27	6
MOTA	423	OG1		53		48.255	25.053	51.641	1.00 29.51	8
MOTA	424	CG2		53		49.522	27.076	51.502	1.00 24.66	6
ATOM ATOM	425 426	С 0	THR A	53 53		46.908 45.778	25.462 25.228	49.209 49.634	1.00 26.58 1.00 21.98	6
MOTA	427	Ŋ	LYS A	54		47.455	24.782	48.203	1.00 21.98	8 7
MOTA	428	CA	LYS A	54		46.739	23.713	47.507	1.00 32.62	6
ATOM	429	CB	LYS A	54		47.601	23.151	46.370	1.00 31.99	6
ATOM	430	CG	LYS A	54		46.985	21.967	45.629	1.00 36.62	6
MOTA	431	CD	LYS A	54		45.733	22.352	44.866	1.00 40.69	6
MOTA	432	CE	LYS A	54		46.058	23.173	43.625	1.00 46.44	6
MOTA	433 434	NZ C	LYS A	54 54		46.844 46.348	22.393 22.595	42.614 48.465	1.00 50.68	7 6
MOTA MOTA	435	ō	LYS A	54		45.277	21.991	48.330	1.00 36.00 1.00 34.77	8
MOTA	436	N	GLU A	55		47.216	22.336	49.443	1.00 37.91	7
MOTA	437	CA	GLU A	55		46.979	21.290	50.433	1.00 36.96	6
MOTA	438	CB	GLU A	55		48.240	21.100	51.281	1.00 40.29	6
ATOM	439	CG	GLU A	55		48.216	19.887	52.195	1.00 47.95	6
MOTA	440	CD	GLU A	55		49.552	19.654	52.891	1.00 51.01	6
ATOM	441		GLU A	55		49.659	18.688	53.679	1.00 52.65	8
ATOM	442 443	OE2 C	GLU A GLU A	55 55		50,497 45.771	20.437 21.609	52.646 51.322	1.00 51.27 1.00 34.10	8 6
MOTA -	444	Ö	GLU A	55	•	44.892	20.769	51.322	1.00 34.10	8
ATOM	445	Ŋ	GLU A	56		45.723	22.827	51.866	1.00 32.39	7
ATOM	446	CA	GLU A	56		44.621	23.256	52.733	1.00 30.13	6
ATOM	447		GLU A	56		44.824	24.714	53.177	1.00 25.28	6
ATOM	448	CG	GLU A	56		46.204	24.994	53.758	1.00 28.82	6
MOTA	449	CD	GLU A	56		46.421	26.450	54.181	1.00 30.74	6
MOTA	450		GLU A	56		46.072	27.369	53.398	1.00 29.77	8
ATOM	451 452	OE2 C	GLU A GLU A	56 56		46.969 43.264	26.674 23.114	55.288 52.024	1.00 25.98 1.00 29.63	8 6
ATOM ATOM	453	0	GLU A	56		42.299	22.584	52.595	1.00 29.90	8
ATOM	454	N	LEU A	57		43.188	23.581	50.780	1.00 26.76	7
MOTA	455	CA	LEU A	57		41.944	23.490	50.020	1.00 25.29	6
ATOM	456	CB	LEU A	57		42.132	24.103	48.629	1.00 22.68	6
ATOM	457	CG	LEU A	57		42.402	25.612	48.572	1.00 22.39	6
ATOM	458		LEU A .			42.654	26.045	47.123	1.00 20.77	. 6
ATOM	459		LEU A	57		41.211	26.366	49.156	1.00 17.66	6
ATOM	460	C	LEU A	57 57		41.479	22.037	49.896	1.00 26.02 1.00 23.41	6 8
	461	И О .	LEU A	57 58		40.284 42.444	21.741 21.143	50.014 49.675	1.00 23.41	7
ATOM	462	IN .	LEU A			46.444	21.14J	49.675	24.02	•

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MOTA	463	CA	LE	ΙL	58	42.194	1 19.718	49.526	1.00 22.44	6
ATOM	464	CB	LĘ	JA	58	43.434	19.027	48.965	1.00 21.02	6
ATOM	465	CG	LE	JA	58	43.838		47.558	1.00 22.94	6
ATOM	466	CD	l LE	JA	58	45.212		47.176	1.00 20.35	6
ATOM	467	CD	2 LET	JA	58	42.755		46.587	1.00 23.28	6
ATOM	468	С	LEU	JΑ	58	41.797		50.835	1.00 25.20	6
MOTA	469		LE		58	41.456		50.854	1.00 26.55	8
ATOM	470		LEU		59	41.858		51.938	1.00 25.44	7
ATOM	471		LEU		59	41.446		53.211	1.00 25.24	
ATOM	472		LEU		59	41.559				6
ATOM	473	CG	LEU		59	42.956		54.350	1.00 24.68	6
MOTA	474		LEU		59	42.912		54.912	1.00 27.05	6
ATOM	475		LEU		59	43.492		56.001	1.00 24.76	- 6
ATOM .	. 476		LEU		59			55.474	1.00 26.99	6
	477	Ö	LEU			39.991		53.045	1.00 24.22	6
MOTA	478				59	39.548		53.581	1.00 21.18	8
ATOM		N	PHE		60	39.270		52.270	1.00 25.00	7
MOTA	479	CA	PHE		60	37.859		52.011	1.00 25.00	6
ATOM	480	CB	PHE		60	37.054		52.605	1.00 26.34	6
MOTA	481	CG	PHE		60	35.600		52.223	1.00 29.37	6
MOTA	482		PHE		60	34.811		52.427	1.00 27.57	6
ATOM	483		PHE		60	35.015		51.661	1.00 27.33	6
MOTA	484		PHE		60	33.466		52.077	1.00 27.00	6
MOTA	485	CE2			60	33.670		51.306	1.00 28.08	6
ATOM	486	CZ	PHE		60	32.893		51.513	1.00 29.48	6
MOTA	487	C	PHE		60	37.506		50.538	1.00 27.78	6
ATOM	488	0	PHE		60	37.022		50.158	1.00 31.57	. 8
ATOM	489	N	HIS		61	37.734		49.696	1.00 26.76	7
ATOM	490	CA	HIS		61	37.376		48.287	1.00 28.84	6
ATOM	491	CB	HIS		61	37.365		47.561	1.00 27.76	6
MOTA	492	CG	HIS		61	36.385		48.117	1.00 30.54	6 '
MOTA	493		HIS		61	35.056	22.549	47.907	1.00 33.74	6
MOTA	494		HIS		61	36.750	23.401	48.987	1.00 34.02	7
MOTA	495		HIS		61.	35.691	24.135	49.286	1.00 32.07	6
MOTA	496	NE2			61	34.649	23.638	48.644	1.00 34.10	7
MOTA	497	С	HIS	A	61	38.278	19.056	47.539	1.00 28.38	6
ATOM	498	0	HIS	Α	61	39.287	18.604	48.072	1.00 25.81	8
ATOM	499	N	THR		62	37.895	18.705	46.310	1.00 32.88	7
ATOM	500	CA	THR		62	38.658	17.749	45.488	1.00 34.68	6
MOTA	501	CB	THR	Α	62	37.715	16.739	44.778	1.00 34.36	6
MOTA	502	0G1	THR	Α	62	36.942	17.415	43.778	1.00 34.81	8
ATOM	503	CG2	THR	A	62	36.759	16.112	45.778	1.00 34.33	6
ATOM	504	С	THR	Α	62	39.485	18.454	44.408	1.00 35.60	6
ATOM	505	0	THR	A	62	39.017	19.418	43.790	1.00 30.85	8
ATOM	506	N	GLU	А	63	40.700	17.958	44.166	1.00 37.38	7
ATOM	507	CA	GLU	A	63	41.587	18.555	43.165	1.00 40.68	6
ATOM	508	CB	GLU	A	63	42.759	17.626	42.840	1.00 43.75	6
MOTA	509	CG	GLU	A	63	43.719	17.389	43.987	1.00 50.68	6
ATOM	510	CD	GLU	A	63	45.026	16.760	43.529	1.00 55.36	6
ATOM	511	OE1	GLU	A	63	45.789	17.441	42.808	1.00 53.03	8
MOTA	512	OE2	GLU	A	63	45.285	15.585	43.883	1.00 59.56	8
ATOM	513	С	GLU	A	63	40.894	18.939	41.860	1.00 39.26	6
ATOM	514	0	GLU	Α	63	40.771	20.116	41.535	.1.00 42.33	8
ATOM	515	N	ASP		64	40.453	17.948	41.102	1.00 37.07	7
ATOM	516	CA	ASP		64	39.782	18.224	39.845	1.00 36.98	6
ATOM	517	CB	ASP		64	38.957	17.000	39.426	1.00 42.19	6
ATOM	518	CG	ASP		64	38.037	16.501	40.533	1.00 47.66	6
MOTA	519		ASP		64	37.039	17.193	40.851	1.00 47.95	8
ATOM	520		ASP		64	38.325	15.413	41.091	1.00 47.95	·8
ATOM	521	c	ASP		64	38.908	19.480	39.906		0
	522	0	ASP		64				1.00 33.40	6
ATOM	523	N	TYR .		65	38.927	20.293	38.986	1.00 33.64	8
ATOM	524	CA	TYR		65	38.156 37.286	19.641	40.990	1.00 30.57	7
ATOM	525	CB	TYR		65		20.806	41.157	1.00 29.65	6
MOTA	526	CG				36.300	20.560	42.316	1.00 30.16	6
ATOM			TYR .		65	35.557	21.790	42.810	1.00 28.49	6
ATOM	527				65	34.791	22.572	41.944	1.00 30.25	6.
ATOM	528	CEL	TYR .	A	65	34.126	23.715	42.399	1.00 28.36	6

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MOTA	529		2 TY	R A	65	35	. 638	22.181	44.150	1.00	28.28	6
ATOM	530		_			34	.980	23.320	44.617		26.96	6
ATOM	531			R A			.227	24.082	43.740	1.00	29.79	6
MOTA	532			RA			.568	25.201	44.207	1.00		8
MOTA	533 534			R A	65		.118	22.061	41.418	1.00		6
MOTA MOTA	535			R A E A	65		.860	23.128	40.857		30.45	8
ATOM	536			EA	66 66		.122	21.926	42.270		26.61	7
MOTA	537			EA	66		.986	23.041	42.597	1.00		6
ATOM	538				66		.009	22.652 23.753	43.687	1.00		6
MOTA	539		1 ILI		66		.264	22.341	43.869 44.992	1.00 1.00		6
ATOM	540		1 IL		66		.478	23.517	45.555	1.00		6 6
ATOM	541	С	ILE	E A	66		.761	23.504	41.381	1.00		6
ATOM	542	0	ILE	A 3	66		. 039	24.696	41.225	1.00		8
ATOM	543	N	ASN		67	41	.125	22.559	40.521	1.00		7
ATOM	544	CA	ASN		67	. 41	.902	22.898	39.337	1.00		6
ATOM	545	CB	ASN		67		.563	21.656	38.726	1.00	34.20	6
MOTA	546	CG	ASN		67		.712	21.118	39.578	1.00		6
MOTA	547 548		1 ASN		67		. 674	21.841	39.878	1.00		8
ATOM ATOM	549	ND:	2 ASN ASN		67 67		.626	19.845	39.956	1.00		7
MOTA	550	ō	ASN		67		.020 .494	23.554 24.354	38.314	1.00		6
ATOM	551	N	THR		68		733	23.221	37.499 38.361	1.00		8 7
ATOM	552	CA	THR		68		787	23.791	37.416	1.00		6
MOTA	553	CB	THR	Α	68		438	23.111	37.500	1.00		6
MOTA	554	OG1			68	37.	620	21.695	37.371	1.00		8
ATOM	555	CG2			68	36.	549	23.591	36.359	1.00	17.59	6
ATOM	556	C	THR		68		633	25.263	37.732	1.00		6
MOTA	557	0	THR		68		529	26.088	36.830	1.00		8
ATOM ATOM	558 559	N CA	LEU		69 69		645	25.582	39.023	1.00		. 7
ATOM	560	CB	LEU		69		535 376	26.956 26.982	39.482	1.00		6
ATOM	561	CG	LEU		69		023	26.527	41.000 41.548	1.00 2		6 6
MOTA	562	CD1			69		087	26.416	43.066	1.00		6
ATOM	563	CD2			69		942	27.528	41.120	1.00		6
MOTA	564	С	LEU	Α	69	39.	772	27.757	39.088	1.00 2		6
ATOM	565	0	LEU		69		683	28.921	38.674	1.00 2		8
ATOM	566	N	MET		70		932	27.128	39.218	1.00 2		7
MOTA MOTA	567 568	CA CB	MET MET		70		183	27.794	38.897	1.00 2		6
ATOM	569	CG	MET		70 70		358 418	26.953 26.751	39.380	1.00 2		6
ATOM	570	SD	MET		70		970	25.929	40.884 41.325	1.00 2		6 16
ATOM	571	CE	MET		70		137	27.077	40.642	1.00 2		6
ATOM	572	C	MET	Α	70		324	28.040	37.412	1.00 2		6
ATOM	573	0	MET	Α	70	42.	903	29.041	36.982		8.99	8
MOTA	574	N	GLU		71	41.		27.122	3 <i>६</i> 632	1.00 2		7
ATOM	575	CA	GLU		71	41.		27.204	31.189	1.00 2	4.41	6
ATOM	576	CB	GLU		71	41.		25.814	34 582	1.00 2		6
ATOM	577 578	CG CD	GLU		71	42.		25.695	33.167	1.00 3		6
ATOM ATOM	579	OE1	GLU		71 71	43. 44.		25.905 25.855	33.099	1.00 3		6
ATOM	580		GLU		71	44.		26.116	31.983 34.154	1.00 3		8
ATOM	581	c	GLU		71	40.		28.160	34.592	1.00 2		8 6
ATOM	582	0	GLU		71	41.		28.851	33.626	1.00 2		8
ATOM	583	N	ALA		72	39.		28.197	35.169	1.00 1		7
ATOM	584	CA	ALA	A	72	38.	589	29.067	34.684	1.00 1		6
ATOM	585	CB	ALA		72	37.		28.743	35.397	1.00 1		6
MOTA	586	C	ALA		72	38.		30.536	34.899	1.00 2		6
ATOM	587	0	ALA		72	38.		31.383	34.016	1.00 2		8
MOTA	588	N	GLU		73	39.4		30.835	36.079	1.00 2		7
MOTA MOTA	589 590	CA CB	GLU		73 73	39.1		32.202	36.436	1.00 2		6
ATOM	591	CG	GLU		73	40.1 40.1		32.282 33.655	37.931 38.349	1.00 2		6
MOTA	592	CD	GLU		73	40.8		33.806	39.841	1.00 2		6 6
ATOM	593.		GLU		73	39.8		33.776	40.582	1.00 3		. 8
ATOM	594		GLU		73	41.9		33.960	40.277	1.00 3		8
									-			

ATOM	595	С	GLU A	A 73	40.946	32.840	35.615	1.00 31.83	6
ATOM	596	ō	GLU Z		40.859	34.024	35.259	1.00 33.52	8
			_						
MOTA	597	N	ARG A		41.992	32.071	35.309	1.00 31.45	7
ATOM	598	CA	ARG A	A 74	43.128	32.611	34.557	1.00 30.65	6
MOTA	599	CB	ARG A	A 74	44.405	31.826	34.874	1.00 32.12	6
ATOM	600	CG	ARG A		44.514	30.467	34.205	1.00 31.42	6
MOTA	601	CD	_ARG A		45.702	29.714	34.754	1.00 30.73	6
MOTA	602	NE	ARG A	4 74	46.041	28.561	33.933	1.00 34.18	7
ATOM	603	CZ	ARG A	A 74	46.646	28.634	32.748	1.00 35.55	6
	604		1 ARG A		46.989		32.232		7
ATOM						29.818		1.00 29.64	
MOTA	605	NH:			- 46.906	27.514	32.079	1.00 34.07	7
ATOM	606	С	ARG A	74	42.894	32.623	33.051	1.00 28.61	6
ATOM	607	0	ARG A	4 74	43.431	33.465	32.338	1.00 24.38	8
MOTA	608	N	CYS A		42.107	31.673	32.566	1.00 28.32	7
MOTA	609	CA	CYS ?		41.796	31.619	31.148	1.00 32.42	5
MOTA	610	CB	CYS. A	¥ 75	41.687	30.167	30.682	1.00 32.91	6
MOTA	611	SG	CYS A	75	43.281	29.296	30.777	1.00 37.09	16
ATOM	612	С	CYS A		40.489	32,382	30.956	1.00 33.10	6
MOTA	613	0	CYS A		40.029	32.598	29.834	1.00 30.74	8
ATOM	614	N	GLN A	3 76	39.914	32.787	32.088	1.00 34.42	7
ATOM	61,5	CA	GLN A	76	38.691	33.575	32.144	1.00 33.20	6
ATOM	616	CB	GLN A	76	38.986	34.962	31.578	1.00 32.09	5
	617	CG	GLN A		38.089				
ATOM						36.064	32.094	1.00 39.46	6
ATOM	618	CD	GLN A		38.479	36.541	33.480	1.00 41.47	્6
ATOM	619	OE:	L GLN 🤉	76	38.574	35.755	34.426	1.00 45.02	8
ATOM	620	NE2	GLN A	76	38.703	37.846	33.606	1.00 42.22	7
ATOM	621	С	GLN A		37.561	32.920	31.358	1.00 33.20	6
MOTA	622	0	GLN A		. 36.732	33.598	30.760	1.00 34.19	8
MOTA	623	N	CYS A		37.522	31.598	31.370	1.00 31.81	7
ATOM	624	CA	CYS A	77	36.511	30.862	30.627	1.00 31.47	б
MOTA	625	CB	CYS . A	. 77	37.187	30.181	29.454	1.00 30.25	6
ATOM	626	SG	CYS A		38.479	29.071	30.044		
								1.00 33.94	16
MOTA	627	С	CYS A		35.851	29.795	31.498	1.00 31.97	6
MOTA	628	Ö	CYS A	. 77	36.335	29.503	32.590	1.00 35.15	8
ATOM	629	N	VAL A	78	34.750	29.216	31.018	1.00 30.78	7
ATOM	630	CA	VAL A		34.069	28.139	31.747	1.00 30.55	6
		CB	VAL A						
ATOM	631				32.539	28.287	31.720	1.00 30.06	6
ATOM	632	CG1	. VAL A	. 78	31.881	27.030	32.293	1.00 28.23	б
ATOM	633	CG2	VAL A	. 78	32.129	29.503	32.526	1.00 30.67	6
ATOM	634	С	VAL A	. 78	34.420	26.794	31.110	1.00 29.80	6
ATOM	635	0	VAL A		33.851	26.422	30.077	1.00 29.65	8
ATOM	636	N	PRO A		35.337	26.033	31.739	1.00 28.55	7
MOTA	637	CD	PRO A	. 79	35.985	26.335	33.025	1.00 24.39	6
ATOM	638	CA	PRO A	. 79	35.793	24.724	31.261	1.00 28.89	6
ATOM	639	CB	PRO A	. 79	36.622	24.218	32.434	1.00 24.49	6
ATOM	6.0	CG	PRO A		37.239	25.500	32.922	1.00 25.68	6
MOTA	61	С	PRO A		34.668	23.776	30.881	1.00 30.13	6
MOTA	6-2	0	PRO A	. 79	33.697	23.624	31.615	1.00 30.87	8
MOTA	643	N	LYS A	80	34.796	23.136	29.727	1.00 33.44	7
ATOM	644	CA	LYS A	80	33.758	22.216	29.303	1.00 38.52	6
	645	CB	LYS A		34.202		28.076	1.00 45.18	6
MOTA	.					21.421			_
MOTA	646	CG	LYS A		35.450	20.589	28.278	1.00 55.18	6
ATOM	647	CD	LYS A	80	35.788	19.827	27.000	1.00 60.80	6
ATOM	. 648	CE	LYS A	80	37.035	18.976	27.168	1.00 64.25	6
ATOM	649	NZ	LYS A		37.367	18.252	25.911	1.00 68.95	7
ATOM	650	С	LYS A		33.411	21.267	30.443	1.00 36.56	6
ATOM	651	0	LYS A	80	34.293	20.775	31.164	1.00 31.61	8
ATOM	652	N	GLY A	81	32.112	21.035	30.602	1.00 32.57	7
ATOM	653	CA	GLY A	81	31,634	20.155	31.648	1.00 29.81	6
					31.477				6
ATOM	654	C	GLY A	81		20.884	32.965	1.00 28.30	
ATOM	655	0	GLY A	81	30.544	20.612	33.723	1.00 25.49	8
ATOM	656	N	ALA A	82	32.380	21.830	33.218	1.00 25.99	7
ATOM	657	CA	ALA A	82	32.384	22.602	34.458	1.00 26.72	6
	658	CB	ALA A	82	33.485	23.674	34.406	1.00 22.64	6
ATOM					_				
ATOM	659	С	ALA A	82	31.066	23.245	34.886	1.00 27.84	6
ATOM	660	0	ALA A	82	30.729	23.224	36.068	1.00 30.00	8

MOTA MOTA MOTA MOTA MOTA	661 662 663 664 665	CA	AR AR AR	G A G A G A	83 83 83		30.310 29.071 28.285 27.439	24.462 24.941 26.189	34.345 33.127 33.408	1.00 32.50 1.00 37.19 1.00 42.23	7 6 6
ATOM-	666	NE		3 A			26.480 25.904			1.00 48.02	6
ATOM	667	CZ		S A			25.046			1.00 53.00	7
ATOM	668		1 ARC				24.649			1.00 56.84	6 7
MOTS	669		2 ARG				24.588			1.00 58.03	7
ATOM	670	С	-AR(3 A	83		28.208		35.189	1.00 31.50	· 6
MOTA	671	0		3 A	83		28.056	23.749	36.386	1.00 29.62	8
MOTA	672	N		JA	84		27.648		34.581	1.00 33.06	7
ATOM	673	CA		JA	94	•	26.819		35.343	1.00 35.40	6
ATOM ATOM	· 674 675	CB CG	GLU GLU		84 84		26.112		34.417	1.00 37.35	· 6
ATOM	676	CD	GLU		84		26.989 27.551		33.496 32.267	1.00 40.01	6
ATOM	677		1 GLU		84		27.925		31.292	1.00 44.49 1.00 41.12	6 -
MOTA	678		2 GLU		84		27.636		32.270	1.00 41.12	8
MOTA	679	С	GLU	J A	84		27.617		36.417	1.00 35.42	6
ATOM	680	0	GLU		84		27.246	20.816	37.594	1.00 34.66	8
ATOM	681	N	LYS		85		28.727	20.226	36.002	1.00 35.21	7
ATOM ATOM	682 683	CA	LYS		85		29.604	19.450	36.878	1.00 37.93	6
ATOM	684	CB	LYS		85 85		30.841 31.739	19.030 17.977	36.076	1.00 40.61	6
·ATOM	685	CD	LYS		85		31.739	16.640	36.706 36.872	1.00 42.63	6
ATOM	686	CE	LYS		85		32.054	15.523	37.078	1.00 45.48	6 6
atom	687	NZ	LYS	A	85		33.032	15.833	38.154	1.00 46.16	7
ATOM	688	С	LYS		85		30.032	20.159	38.175	1.00 37.56	6
ATOM	689	0	LYS		85		30.161	19.516	39.222	1.00 38.40	8
MOTA	690	N	TYR		86		30.254	21.472	38.116	1.00 35.60	7
MOTA MOTA	691 692	CA CB	TYR TYR		86 86		30.671	22.216	39.307	1.00 32.67	6
ATOM	693	CG	TYR		86		32.151 33.065	22.610 21.424	39.200	1.00 32.09	6
MOTA	694		TYR		86		33.120	20.393	38.995 39.932	1.00 33.63 1.00 32.12	ნ 6
ATOM	695		TYR		86		33.918	19.266	39.723	1.00 32.12	. 6
MOTA	696	CD2	TYR	Α	86		33.839	21.306	37.841	1.00 33.82	6
ATOM	697		TYR		86		34.645	20.178	37.623	1.00 34.55	6
ATOM	698	CZ	TYR		86		34.675	19.162	38.566	1.00 32.38	6
ATOM ATOM	699 700	OH C	TYR TYR		86 86		35.431	18.034	38.336	1.00 29.17	8
ATOM	701	ō	TYR		86		29.831 30.192	23.455 24.265	39.597 40.445	1.00 30.21 1.00 29.12	6
ATOM	702	N	ASN		87		28.712	23.594	38.893	1.00 29.12	8 7
MOTA	703	CA	ASN	A	87		27.797	24.717	39.086	1.00 28.58	6
ATOM	704	CB	ASN	Α	87		27.154	24.618	40.470	1.00 25.63	6
ATOM	705	CG	ASN		87		25.871	25.428	40.596	1.00 28.05	6
ATOM	706 707		ASN		87		25.275	25.477	41.672	1.00 71.32	8
ATOM ATOM	707	C MDS	ASN ASN		87 87		25.434 28.580	26.055	39.506	1.00 .8.35	7
ATOM	709	õ	ASN		87		28.319	26.015 26.981	38.963 39.677	1.00 10.35	6 8
ATOM	710	N	ILE		88		29.545	26.019	38.051	1.00 32.07	. 5
ATOM	711	CA	ILE		88		30.407	27.173	37.809	1.00 33.77	6
ATOM	712	CB	ILE		88		31.894	26.734	37.776	1.00 36.13	6
ATOM	713		ILE		88		32.759	27.831	37.201	1.00 37.80	6
ATOM	714	CG1	ILE	À	88		32.357	26.342	39.178	1.00 38.92	6
ATOM	715 716	CDI	ILE		88		32.350	27.483	40.176	1.00 41.44	6
ATOM ATOM	717	0	ILE		88 88		30.085 29.708	27.857 27.196	36.482	1.00 32.28 1.00 32.72	6
ATOM	718	N	GLY	A	89		30.237	29.179	35.520 36.438	1.00 32.72	8 7
ATOM	719	CA	GLY	A	89		29.994	29.915	35.207	1.00 30.84	6
ATOM	720	С	GLY		89		28.696	30.689	35.093	1.00 32.17	6
ATOM	721	0	GLY		89		28.628	31.670	34.349	1.00 30.42	8
ATOM	722	N	GLY		90		27,670	30.257	35.821.	1.00 31.51	7
ATOM	723	CÀ	GLY		90		26.387	30.937	35.756	1.00 32.92	5
ATOM ·	724 725		GLY GLY		90 90		26.316 27.302	32.246 32.671	36.524 37.129	1.00 34.32 1.00 33.97	6 8
ATOM		N	TYR .	λ	91		25.144	32.871	36.504	1.00 33.97	7
••••								J2.002		2.00	•

ATOM	727	CA	TYR A	91	24.924	34.146	37.206	1 00 35 76	_
								1.00 35.76	6
ATOM	728		TYR A		23.465	34.589	37.058	1.00 38.10	6
ATOM	729	CG	TYR A	91	23.089	35.733	37.990	1.00 42.40	6
MOTA	730	CD1	L TYR A	91	23.417	37.057	37.688	1.00 43.57	6
MOTA	731				23.105				
						38.106	38.577	1.00 42.44	5
MOTA	732		TYR A	91	22.444	35.484	39.205	1.00 44.07	6
ATOM	733	CE2	TYR A	91	22.132	36.526	40.097	1.00 42.67	6
MOTA	734	CZ	TYR A	91	22.462	37.825	39.775	1.00 42.41	
	735								6
ATOM			TYR A		22.130	38.835	40.646	1.00 43.69	8
MOTA	736	С	TYR A	. 91	25.242	34.082	38.701	1.00 34.15	6
MOTA	737	0	TYR A	91	25.821	35.014	39.266	1.00 29.52	8
ATOM	738	N	GLU A	92	24.837	32.986	39.333	1.00 34.78	7
ATOM	. 739		GLU A		25.024				
	_					32.797	40.767	1.00 38.46	6
MOTA	740		GLU A		24.233	31,564	41.211	1.00 43.99	6
ATOM	741	CĠ	GLU A	92	23.932	31.489	42.700	1.00 52.10	. 6
ATOM	742	CD	GLU A	92	23.294	30.161	43.097	1.00 58.00	. 6
MOTA	743		GLU A		24.001	29.126	43.058		
								1.00 60.63	8
ATOM	744	OE2			22.087	30.149	43.434	1.00 59.58	8
ATOM	745	С	GLU A	92	26.492	32.669	41.208	1.00 36.42	6
ATOM	746	0	GLU A	92	26.902	33.287	42.193	1.00 32.92	8
ATOM	747	N	ASN A	93	27.280	31.883	40.473	1.00 34.12	7
ATOM	748	CA	ASN A		28.693				
						31.671	40.808	1.00 33.24	6
ATOM	749	CB	ASN A	93	28.871	30.259	41.364	1.00 28.52	6
ATOM .	750	CG	ASN A	93	27.734	29.859	42.299	1.00 27.45	6
MOTA	751	OD1	ASN A	93	27.547	30.457	43.355	1.00 21.76	8
ATOM	752		ASN A	93	26.956	. 28.853	41.895		7
								1.00 21.79	
MOTA	753	C	ASN A	93	29.529	31.843	39.535	1.00 35.04	6
MOTA	754	0	ASN A	93	30.160	30.898	39.059	1.00 33.81	8
ATOM	755	N	PRO A	94	29.583	33.081	39.010	1.00 36.19	7
ATOM	756	CD	PRO A	94	28.970	34.231	39.690	1.00 34.62	6
MOTA	757	CA	PRO A	94	30.274	33.560			
							37.808	1.00 34.80	6
ATOM	758	CB	PRO A	94	29.924	35.050	37.791	1.00 33.94	6
ATOM	759	CG	PRO A	94	28.619	35.095	38.516	1.00 36.13	6
MOTA	760	С	PRO A	94	. 31.775	33.379	37.733	1.00 34.63	6
ATOM	761	Ō	PRO A	94	32.443	33.103	38.730		
								1.00 34.72	8
ATOM	762	N	VAL A	95	32.299	33.556	36.526	1.00 33.57	7
MOTA	763	CA	VAL A	95	33.735	33.499	36.307	1.00 30.31	6
ATOM	764	CB	VAL A	95	34.085	33.171	34.841	1.00 29.88	6
ATOM	765	CG1	VAL A		35.561	33.453	34.574	1.00 29.53	6
ATOM	766		VAL A	95	33.795				
						31.713	34.563	1.00 28.05	6
ATOM	767	С	VAL A	95	34.195	34.910	36.624	1.00 29.86	6
ATOM	768	0	VAL A	95	33.524	35.879	36.272	1.00 29.07	8
ATOM	769	N	SER A	96	35.318	35.019	37.317	1.00 30.89	7
ATOM	770	CA	SER A	96	35.889	36.310	37.687	1.00 32.27	6
ATOM	771	СВ	SER A	96	34.885	37.145	38.501		
								1.00 30.16	6
MOTA	772	OG	SER A	96	34.600	36.545	39.756	1.00 26.77	8
MOTA	773	С	SER A	96	37.111	35.993	38.537	1.00 32.96	6
MOTA	774	0	SER A	96	37.603	34.865	38.511	1.00 33.77	8
ATOM	775	N	TYR A	97	37.609	36.973	39.282	1.00 32.66	7
ATOM	776	CA	TYR A	97	38.753	36.712	40.132	1.00 31.95	6
MOTA	777	CB	TYR A	97	39.838	37.766	39.923	1.00 31.81	6
ATOM	778	CG	TYR A	97	40.416	37.729	38.525	1.00 30.39	6
MOTA	779	CD1	TYR A	97	39.820	38.434	37.479	1.00 30.63	6
ATOM	780		TYR A	97	40.327	38.358	36.178	1.00 28.49	6
ATOM	781		TYR A	97	41.536	36.945	38.236	1.00 28.43	6
MOTA	782	CE2	TYR A	97	42.046	36.858	36.942	1.00 24.73	6
MOTA	783	CZ	TYR A	97	41.437	37.565	35.919	1.00 27.27	6
ATOM	784	OH	TYR A	97	41.915	37.455	34.633	1.00 26.70	8
	785		TYR A	97	38.350	36.618	41.596	1.00 31.10	6
MOTA									
ATOM	786		TYR A	97	39.178	36.735	42.495	1.00 33.01	. 8
ATOM	787		ALA A	98	37.059	36.398	41.818	1.00 31.11	. 7
ATOM	738	CA	ALA A	98	36.510	36.241	43.160	1.00 30.06	5
ATOM	789		ALA A	98	35.141	36.920	43.256	1.00 27.71	6
			ALA A						
ATCM	790	-		98	36.350	34.736	43.357	1.00 31.24	. 6
ATCM	791		ALA A	98	36.335	34.238	44.487	1.00 29.66	-8
ATOM	792	N	MET A	99	36.249	34.030	42.230	1.00 29.50	7
				•			-		
					•				

MOTA	793 CA MET A 99	36.04	0 33 50			
MOTA	794 CB MET A 99	35.77				
ATOM	795 CG MET A 99					6
ATOM		36.94				6
MOTA		36.42			5 1.00 29.78	16
		35.62		38.34	7 1.00 25.05	- 6
MOTA	798 C MET A 99	37.19		42.783		6
MOTA	799 O MET A 99	36.99	30.757		1.00 30.59	8
MOTA	800 N PHE A 100	38.41				
MOTA	801 CA PHE A 100	39.55				7
MOTA	802 CB PHE A 100	40.322			1.00 33.87	6
MOTA	803 CG PHE A 100	41.43				6
ATOM	804 CD1 PHE A 100	41.152				6
MOTA	805 CD2 PHE A 100					6
ATOM	806 CE1 PHE A 100	42.768				6
ATOM	807 CE2 PHE A 100					6
	808 CZ PHE A 100	43.808			1.00 40.50	6
ATOM		43.517		43.729	1.00 39.89	6
ATOM	809 C PHE A 100	40.519			1.00 33.98	6
ATOM	810 O PHE A 100	40.706	32.231	45.088	1.00 38.21	8
ATOM	811 N THR A 101	41.137	33.415		1.00 28.09	
MOTA	812 CA THR A 101	42.063	34.261		1.00 22.19	7
MOTA	813 CB THR A 101	42.623	35.378	43.072		6
ATOM	814 OG1 THR A 101	43.441	34.795	42.052	1.00 22.48	6
ATOM	815 CG2 THR A 101	43.468				8
ATOM	816 C THR A 101	41.408	36.335	43.876		6
ATOM	817 O THR A 101	41.988		45.205		6
ATOM	818 N GLY A 102		34.845	46.282	1.00 23.82	8
ATOM	819 CA GLY A 102	40.197	35.377	45.068	1.00 21.79	7
ATOM		39.533	35.947	46.231	1.00 21.23	6
ATOM		39.072	34.833	47.153	1.00 23.03	6
		39.209	34.909	48.378	1.00 20.41	8
ATOM	822 N SER A 103	38.512	33.792	46.544	1.00 22.59	7
ATOM	823 CA SER A 103	38.028	32.640	47.276	1.00 26.51	6
ATOM	824 CB SER A 103	37.454	31.598		1.00 28.10	6
ATOM	825 OG SER A 103	36.314	32.099		1.00 32.01	
MOTA	826 C SER A 103	39.188	32.040	48.032	1.00 32.01	8
ATOM	827 O SER A 103	39.019	31.544	49.144	1.00 27.73	6
ATOM	828 N SER A 104	40.364	32.080		1.00 30.61	8
ATOM	829 CA SER A 104	41.590		47.410	1.00 28.76	7
ATOM	830 CB SER A 104	42.769	31.552	48.008	1.00 28.55	6
ATOM	831 OG SER A 104		31.683	47.039	1.00 28.74	6
ATOM	832 C SER A 104	42.501	31.044	45.804	1.00 35.04	8
MOTA	833 O SER A 104	41.870	32.401	49.226	1.00 25.67	6
ATOM		42.026	31.897	50.338	1.00 25.17	8
MOTA		41.909	33.705	48.986	1.00 23.91	7
		42.163	34.698	50.008	1.00 23.01 .	6
MOTA	836 CB LEU A 105	42.049	36.082	49.382	1.00 23.57	6
MOTA	837 CG LEU A 105	43.158	37.091	49.672	1.00 26.30	6
MOTA	838 CD1 LEU A 105	44.502	36.551	49.178	1.00 22.38	6
MOTA	839 CD2 LEU A 105	42.823	38.413	48.984	1.00 27.36	6
ATCM	840 C LEU A 105	41.187	34.559	51.182		
MOTA	841 O LEU A 105	41.604	34.448	52.331		. 6
MOTA	842 N ALA A 106	39.887	34.556	50.897	1.00 21.60	8
ATOM	843 CA ALA A 106	38.884			1.00 25.32	7
ATOM	844 CB ALA A 106	20.004	34.423	51.957	1.00 26.04	6
ATOM	845 C ALA A 106	37.471	34.423	51.358	.1.00 24.28	6
ATOM.		39.088	33.158	52.790	1.00 25.76	6
		38.953	33.186	54.015	1.00 22.75	8
MOTA	847 N THR A 107	39.410	32.057	52.111	1.00 25.65	7
ATOM	848 CA THR A 107	39.620	30.760	52.754	1.00 25.54	6
MOTA	849 CB THR A 107	39.706	29.637	51.713	1.00 21.92	6
MOTA	850 OG1 THR A 107	38.559	29.688	50.868	1.00 26.40	
MOTA	851 CG2 THR A 107	39.742	28.295	52.387	1 00 17 76	8
ATOM	852 C THR A 107	40.901	30.720	53.583	1.00 17.36	6
MOTA	853 O THR A 107	40.906			1.00 28.16	6
ATOM	854 N GLY A 108	41.994	30.254	54.727	1.00 28.07	8
ATOM	855 CA GLY A 108		31.191	52.996	1.00 28.51	7
ATOM ·	856 C GLY A 108	43.247	31.187	53.718	1.00 28.37	б
	-	43.027	31.921	55.019	1.00 30.26	6
ATOM ATOM		43.502	31.499	56.076	1.00 32.98	8
ATOM		42.283	33.018	54.942	1.00 24.81	7
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Figure 17-14

ATOM	859	CA	SER A 1	.09	42.002	33.810	56.119	1.00 24.86	6
ATOM	860) CE			41.222		55.727	1.00 24.74	6
ATOM	861	. og			41.992		54.872	1.00 21.07	8
ATOM	862		SER A 1		41.240		57.173	1.00 27.89	6
MOTA	863		SER A 1		41.424		58.377	1.00 30.92	8
ATOM	864		THR A 1		40.389		56.744	1.00 23.91	7
ATOM	865				39.676		57.721	1.00 24.80	6
ATOM	866				38.641		57.074	1.00 29.65	
ATOM	867		1 THR A 1		37.469		56.669		6
ATOM	868		2 THR A 1		38.228		58.067	1.00 30.45	8
ATOM	869		THR A 1		40.712			1.00 29.00	6
ATOM	870		THR A 1		40.615		58.478	1.00 24.34	6
ATOM	871				41.715		59.699	1.00 24.74	8
ATOM	872				42.759		57.764	1.00 23.01	7
ATOM	873		VAL A 1		43.695		58.416	1.00 24.13	6
ATOM	874		1 VAL A 1		44.845		57.391	1.00 25.77	6
ATOM	875		2 VAL A 1		42.888	_	58.121	1.00 22.51	6
ATOM	876		VAL A 1				56.534	1.00 22.67	6
ATOM	877	Ö			43.576		59.329	1.00 23.14	6
ATOM	878	Ŋ	VAL A 1.		43.720		60.518	1.00 24.11	8
ATOM	879	CA	GLN A 1.		44.101	31.156	58.772	1.00 24.94	7
	880				44.895	32.100	59.554	1.00 25.12	6
ATOM	881	CB	GLN A 1		45.082	33.413	58.779	1.00 25.14	6
ATOM ATOM	882	CD	GLN A 1		45.545	33.224	57.330	1.00 28.51	6
ATOM	883		GLN A 1: GLN A 1:		45.789	34.534	56.594	1.00 29.13	6
MOTA	884	NE2			46.779	35.219	56.837	1.00 31.22	8
ATOM	885		GLN A 1.		44.877	34.890	55.694	1.00 29.31	7
ATOM	886	C	GLN A 1.		44.107	32.362	60.827	1.00.24.62	6
ATOM	887	O N	ALA A 1		44.647	32.311	61.939	1.00 21.10	8
	888				42.813	32.622	60.644	1.00 24.41	7
ATOM	889	CA	ALA A 11		41.914	32.904	61.751	1.00 23.33	6
ATOM ATOM	890	CB C	ALA A 11		40.516	33.183	61.224	1.00 19.80	6
ATOM	891	o	ALA A 11		41.901	31.733	62.729	1.00 25.34	6
			ALA A 11		41.925	31.930	63.946	1.00 27.52	8
ATOM	892 893	Й	ILE A 11		41.859	30.509	62.211	1.00 24.39	7
ATOM ATOM	894	CA CB	ILE A 11		41.867	29.356	63.106	1.00 24.49	6
ATOM	895		ILE A 11 ILE A 11		41.524	28.042	62.371	1.00 23.46	6
ATOM	896		ILE A 11		41.902	26.855	63.227	1.00 18.97	6
ATOM	897		. ILE A 11		40.030	28.015	62.034	1.00 21.17	6
ATOM	898	C	ILE A 11		39.598 43.230	26.791	61.239	1.00 22.51	6
MOTA	899	ò	ILE A 11		43.328	29.227	63.757	1.00 24.32	6
ATOM	900	N	GLU A 11			28.817	64.907	1.00 24.74	8
ATOM	901	CA			44.280	29.580	63.019	1.00 26.58	7
ATOM	902	CB	GLU A 11 GLU A 11		45.638	29.518	63.551	1.00 25.89	6
ATOM	903	CG	GLU A 11		46.639	29.992	62.508	1.00 22.63	6
	904	CD			46.554	29.264	61.192	1.00 20.39	6
ATOM ATOM	905		GLU A 11		17.668	29.670	60.244	1.00 21.39	6
	906	OE2	GLU A 11 GLU A 11		47.848	30.887	60.016	1.00 19.60	8
ATOM	907		GLU A 11		18.362	28.769	59.722	1.00 22.53	8
ATOM	908	C	_		45.724	30.422	64.774	1.00 27.56	6
ATOM	909	N O	GLU A 11 GLU A 11		46.173	30.006	65.837	1.00 25.98	8
ATOM					45.267	31.660	64.615	1.00 31.19	7
MOTA	910		GLU A 11		45.282	32.631	65.705	1.00 35.80	6
ATOM	911	CB	GLU A 11		44.676	33.959	65.237	1.00 36.91	6
MOTA	912	CG	GLU A 11		45.434	34.605		1.00 41.14	6
ATOM	913	CD	GLU A 11		46.872	34.982	64.420	1.00 43.09	6
MOTA	914		GLU A 11		47.072	35.886	65.267	1.00 43.42	8
ATOM	915		GLU A 11		47.802	34.369	63.849	1.00 41.76	8
ATOM	916	C	GLU A 11		44.543	32.131	66.947	1.00 35.11	6
MOTA	917	0	GLU A 11		45.054	32.228	68.061	1.00 37.26	8
ATOM	918	N	PHE A 11		43.343	31.598	66.761	1.00 34.30	7
ATOM	919	CA	PHE A 11		42.577	31.096	67.893	1.00 34.44	5
ATOM	920	CB	PHE A 11		41.300	30.399	67.415	1.00 35.45	6
ATOM	921	CG	PHE A 11		40.383	29.979	68.533	1.00 37.14	6
MOTA	922		PHE A 11		39.705	30.930	69.290	1.00 35.80	6
ATCM	923		PHE A 11		40.196	28.630	68.832	1.00 41.05	.6
MOTA	924	CE1	PHE A 11	7	38.853	30.549	70.323	1.00 38:08	·6

SUBSTITUTE SHEET (RULE 26)

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MCTA	925 CE2 PHE A 117	39.33	8 28.234	69.874	1.00 40.44	6
MOTA	926 CZ PHE A 117	38.66	8 29.198			
MOTA	927 C PHE A 117	43.42			1.00 34.24	6
MOTA	928 O PHE A 117	43.49			1.00 34.24	6
MOTA	929 N LEU A 118	44.06				8
ATOM-	930 CA LEU A 118					7
MOTA	931 CB LEU A 118	44.89				6
ATOM		45.15				6
		43.90		67.038	1.00 27.87	6
MOTA	933 CD1 LEU A 118	44.24	4 25.232	65.996	1.00 20.81	6
ATOM	934 CD2 LEU A 118	43.25	9 25.662	68.257		6
MCTA	935 C LEU A 118	46.21	28.696			
MOTA	936 O LEU A 118	46.98				6
MOTA	937 N LYS A 119	46.48		68.843		8
ATOM	938 CA LYS A 119	47.679		69.365		7
MOTA	939 CB LYS A 119	48.143		68.448		6
ATOM	940 CG LYS A 119	48.614				6
MOTA	941 CD LYS A 119		-	67.100		6
MOTA	942 CE LYS A 119	49.111		66.263		6
ATOM	943 NZ LYS A 119	49.691		64.949		6
ATOM		50.167		64.092	1.00 51.48	7
ATOM		47.273		70.705	1.00 34.85	· 6
	945 O LYS A 119	48.112		71.562	1.00 38.12	8
ATOM	946 N GLY A 120	45.967	31.372	70.869	1.00 34.15	7
MOTA	947 CA GLY A 120	45.431	31.927	72.094	1.00 36.25	6
ATOM	948 C GLY A 120	44.860		71.851	1.00 39.02	6
MOTA	949 O GLY A 120	44.640		72.796	1.00 40.23	
ATOM	950 N ASN A 121	44.619		70.586	1.00 38.48	8
ATOM	951 CA ASN A 121	44.079		70.247		7
ATOM	952 CB ASN A 121	44.928	35.624	69.170	1.00 37.47	6
ATOM	953 CG ASN A 121	46.340	35.871		1.00 39.57	6
ATOM	954 OD1 ASN A 121	47.078	34.938	69.622	1.00 41.81	6
ATOM	955 ND2 ASN A 121	46.727		69.926	1.00 47.67	8
ATOM	956 C ASN A 121	42.637	37.134	69.675	1.00 43.63	7
ATOM	957 O ASN A 121		34.893	69.772	1.00 36.59	6
ATOM		42.037	33.818	69.704	1.00 34.08	8
ATOM		42.092	36.061	69.446	1.00 33.53	7
		40.720	36.166	68.976	1.00 34.77	6
ATOM	960 CB VAL A 122	39.861	37.064	69.898	1.00 38.20	6
ATOM	961 CG1 VAL A 122	38.418	37.096	69.388	1.00 37.55	6
ATOM	962 CG2 VAL A 122	39.918	36.553	71.342	1.00 37.77	6
ATOM	963 C VAL A 122	40.731	36.781	67.596	1.00 31.08	6
ATOM	964 O VAL A 122	40.991	37.967	67.441.	1.00 34.19	8
ATOM	965 N ALA A 123	40.451	35.975	66.588	1.00 31.14	7
ATOM	966 CA ALA A 123	40.451	36.476	65.231	1.00 30.26	
ATOM	967 CB ALA A 123	41.307	35.588	64.327	1.00 30.26	6
MCTA	968 C ALA A 123	39.038	36.533	64.716		- 6
ATOM	969 O ALA A 123	38.132	35.924	65.281	1.00 28.26	6
ATOM	970 N PHE A 124	38.875	37.276		1.00 29.28	8
ATOM	971 CA PHE A 124	37.601		63.631	1.00 28.70	7
MCTA	972 CB PHE A 124		37.475.	62.976	1.00 28.38	6
ATOM	973 CG PHE A 124	36.920	38.713	63.563	1.00 29.16	6
ATOM	974 CD1 PHE A 124	35.645	39.099	62.874	1.00 31.20	6
ATOM	975 CD2 PHE A 124	34.679	38.139		1.00 32.00	6
	975 CD2 PRE A 124	35.378	40.435	62.579	1.00 29.53	6
ATOM	976 CE1 PHE A 124	33.463	38.510	61.973	1.00 30.74	6
ATOM	977 CE2 PHE A 124	34.165	40.813	61.988	1.00 27.45	6
ATOM	.978 CZ PHE A 124	33.207	39.847	61.686	1.00 28.72	6
ATOM	979 C PHE A 124	37.880	37.671	61.496	1.00 30.19	6
ATOM	980 O PHE A 124	38.427		61.095	1.00 32.10	
MOTA	981 N ASN A 125	37.545	36.663	60.696	1.00 32.16	8
ATCM	982 CA ASN A 125	37.731	36.728	59.251		7
NOT.	983 CB ASN A 125	38.247	35.393		1.00 30.10	6
TOM	984 CG ASN A 125			58.712	1.00 32.02	6
NCTA	985 OD1 ASN A 125	38.281	35.360	57.195	1.00 33.79	6
	986 ND2 ASN A 125	38.754	36.306	56.556	1.00 31.85	8
TOM		37.790	34.268	56.609	1.00 30.51	7
MOTA	987 C ASN A 125	36.403	37.083	58.584	1.00 29.80	6
MOTA	988 O ASN A 125	35.626	36.206	58.179	1.00 27.24	8
TOM	989 N PRO A 126	36.135	38.386	58.451	1.00 28.25	7
TCM	990 CD PRO A 126	36.997	39.516	58.833	1.00 29.22	6

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1.00 27.92
                                                                            6
                                            38.891
                                                    57.844
                                   34.909
         991
              CA
                  PRO A 126
MOTA
                                                              1.00 29.07
                   PRO A 126
                                   35.139
                                            40.407
                                                     57.856
         992
              CB
MOTA
                                                     57.775
                                                              1.00 26.54
                                   36.649
                                            40.520
         993
              CG
                   PRO A 126
MOTA
                                                     56.448
                                                              1.00
                                                                   27.54
                                   34.651
                                            38.339
         994
                   PRO A 126
              C
MOTA
                                                     55.949
                   PRO A 126
                                    33.532
                                            38.402
                                                              1.00
                                                                   28.66
         995
              O
ATOM
                                                     55.820
                                                              1.00 26.99
                                                                             7
                                   35.687
                                            37.795
MOTA
         996
              N
                   ALA A 127
                                                              1.00 26.54
                                   35.548
                                                     54.477
                                            37.244
         997
              CA
                   ALA A 127
ATOM
                                                                   22.43
                                                                             6
                                    36.822
                                            37.505
                                                     53.684
                                                              1.00
                   ALA A 127
         998
              CB
ÁTOM
                                                     54.480
                                                              1.00 27.38
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                                   35.225
                                            35.744
         999
                   ALA A·127
MOTA
                                                     53.423
                                                              1.00
                                                                   29.04
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                                    35.038
                                            35.140
        1000
              0
                   ALA A 127
MOTA
                                                                             7
                                    35.166
                                            35.142
                                                     55.663
                                                              1.00
                                                                   26.97
                   GLY A 128
MOTA
        1001
              N
                                    34.874
                                            33.724
                                                     55.737
                                                              1.00 25.65
                                                                             6
                   GLY A 128
        1002
              CA
MOTA
                                            33.486
                                                     55.880
                                                              1.00
                                                                   26.17
                                    33.389
                   GLY A 128
        1003
              С
MOTA
                                                                   27.39
                                                              1.00
                                    32.600
                                            34.428
                                                     55.804
                   GLY A 128
        1004
              O
MOTA.
                                                     56.083
                                                             . 1.00 23.87
                                    32.998
                                            32.234
                   GLY A 129
        1005
              N
ATOM
                                                     56.236
                                                                             б
                                            31.936
                                                              1.00
                                                                    25.17
                   GLY A 129
                                    31.588
        1006
              CA
MOTA
                                                     54.937
                                                              1.00
                                                                   25.88
                   GLY A 129
                                    30.847
                                             31.674
              C
        1007
MOTA
                                             31.908
                                                              1.00 25.07
                                                                             8
                                                     54.848
                   GLY A 129
                                    29:643
MOTA
        1008
              0
                                    31.566
                                             31.198
                                                     53.927
                                                              1.00
                                                                   25.69
                                                                             7
                   MET A 130
        1009
              N
MOTA
                                                              1.00 26.48
                                             30.872
                                                     52.622
              CA
                   MET A 130
                                    30.981
        1010
MOTA
                                                     51.567
                                                              1.00 28.53
                                                                             6
                                    32.103
                                             30.907
                   MET A 130
        1011
              CB
ATOM
                                                     51.467
                                                              1.00
                                                                    26.54
                                                                             6
                                    32.795
                                             32.288
        1012
                   MET A 130
              CG
ATOM
                                             32.366
                                                     50.613
                                                              1.00
                                                                    26.29
                                    34.413
                       A 130
        1013
              SD
                   MET
ATOM
                                                     49.062
                                                              1.00 25.85
                                                                             6
                                    34.080
                                             31.512
                   MET A 130
        1014
MOTA
              CE
                                    30.355
                                             29.463
                                                     52.768
                                                              1.00
                                                                    24.47
                                                                             6
        1015
              C
                   MET A 130
MOTA
                                    30.761
                                                     52.113
                                                               1.00 17.67
                                                                             8
                                             28.502
                   MET
                       A 130
        1016
ATOM
                                                     53.636
                                                               1.00
                                                                    23.28
                   HIS A 131
                                             29.389
                                    29.347
        1017
              N
MOTA
                                                               1.00 26.33
                                                                             6
                                                      54.019
                                    28.647
                                             28.161
                   HIS A 131
        1018
              CA
ATOM
                                                     55.180
                                                               1.00 26.98
                                                                             6
                   HIS A 131
                                    27.685
                                             28.485
        1019
              CB
ATOM
                                                               1.00
                                                                    28.50
                                                                             б
                                    26.663
                                             29.540
                                                      54.862
                   HIS A 131
        1020
              CG
MOTA
                                                               1.00 28.65
                                                     53.677
                                    26.225
                                             30.030
                  HIS A 131
ATOM
        1021
              CD2
                                                                             7
                                                      55.831
                                                               1.00 33.04
                                    25.906
                                             30.166
              ND1 HIS A 131
        1022
MOTA
                                                               1.00 27.75
                                             30.995
                                                      55.259
                                                                             6
                                    25.051
        1023
               CE1 HIS A 131
ATOM
                                                      53.952
                                                               1.00 26.97
                                                                             7
                                             30.932
                                    25.224
        1024
              NE2
                   HIS A 131
MOTA
                                                      53.017
                                                               1.00 28.44
                                                                             6
                                             27.284
                                    27.917
                   HIS A 131
ATOM
        1025
              C
                                                                             .3
                                                                    31.15
                                    27.434
                                             26.214
                                                      53,390
                                                               1.00
                   HTS A 131
        1026
              0
ATOM
                                                      51.756
                                                                              7
                                    27.861
                                             27.694
                                                               1.00 30.64
        1027
                   HIS A 132
ATOM
              N
                                                      50.746
                                                                    28.71
                                                                              6
                                             26.938
                                                               1.00
                   HIS A 132
                                    27.111
        1028
              CA
MOTA
                                                               1.00 27.21
                                                      49.890
              CB
                   HIS A 132
                                    26.321
                                             27.941
        1029
MOTA
                                                               1.00 28.83
                                                                              6
                                    25.408
                                             28.819
                                                      50.693
                   HIS A 132
        1030
               CG
 MOTA
                                                      50.578
                                                               1.00
                                                                    28.92
                                                                              6
                                             30.137
                                    25.111
                  HIS A 132
        1031
               CD2
 ATOM
                                                      51.773
                                                               1.00 31.80
                                                                              7
                                             28,360
               ND1 HIS A 132
                                    24.686
        1032
 MOTA
                                                               1.00 29.95
                                                      52.285
               CE1 HIS A 132
                                    23.981
                                             29.353
        1033
 MOTA
                                                                              7
                                                      51.579
                                                               1.00
                                                                    28,21
                                    24.222
                                             30.443
        1034
               NE2
                   HIS A 132
 ATOM
                                                      49.851
                                                               1.00 28.51
                                                                              б
                                    27.889
                                             25.970
        1035
               C
                   HIS A 132
 ATOM
                                                               1.00
                                                                    23.44
                                             24.375
                                                      49.533
                   HIS A 132
                                    27.399.
 MOTA
        1036
               0
                                                               1.00 27.94
                                             26.379
                                                      49.455
                                    29.093
                   ALA A 133
        1037
               N
 MOTA
                                    29.958
                                                      48.579
                                                               1.00 26.99
                                                                              6
                                             25.386
        1038
               CA
                   ALA A 133
 MOTA
                                                      48.392
                                                                              6
                                                               1.00
                                                                     21.87
                                    31.295
                                             26.303
        1039
                   ALA A 133
               CB
 MOTA
                                                               1.00 26.69
                                                                              6
                                                      49.078
                                    30.199
                                             24.164
                   ALA A 133
        1040
               С
 MOTA
                                                                              8
                                             23.973
                                                      50.182
                                                               1.00 28.25
                                    30.703
                   ALA A 133
        1041
               0
 MOTA
                                             23.174
                                                               1.00
                                                      48.255
                                                                     26.73
                                    29.850
                   PHE A 134
        1042
              N
 MOTA
                                                               1.00 25.04
                                                      48.615
                                    30.046
                                             21.773
                   PHE A 134
        1043
               CA
 ATOM
                                                               1.00 19.20
                                                                              6
                                    29.070
                                             20.855
                                                      47.875
                   PHE A 134
        1044
               CB
 MOTA
                                             21.199
                                                               1.00
                                                                     15.75
                                                                              6
                                                      48.100
                                    27.629
                   PHE A 134
               CG
        1045
 ATOM
                                                      47.169
                                                               1.00 14.83
                                    26.929
                                             21.960
                   PHE A 134
               CD1
 MOTA
        1046
                                                               1.00 14.03
                                    26.985
                                             20.814
                                                      49.273
        1047
               CD2
                   PHE
                       A 134
 ATOM
                                             22.336
                                                      47.404
                                                               1.00
                                                                     14.84
                                    25.614
               CE1
                   PHE A 134
        1048
 MOTA
                                                               1.00 12.07
                                                      49.519
                                    25.670
                                              21.184
               CE2 PHE A 134
        1049
 MOTA
                                                               1.00 14.59
                                              21.949
                                                       48.581
                                    24.985
        1050
                   PHE
                       A 134
               CZ
 ATOM
                                             21.310
                                                       48.319
                                                                1.00
                                                                     29.48
                                     31.460
                       A 134
        1051
                   PHE
               C
 ATOM
                                                                1.00 33.19
                                                       47.822
                                     32.291
                                              22.066
                   PHE A 134
         1052
               0
 ATOM
                                                                1.00
                                                                     31.52
                                    31.713
                                              20.045
                                                       48.620
                   LYS A 135
        1053
               N
 ATOM
                                                                1.00
                                                                     29.15
                                     33.012
                                              19.427
                                                       48.427
                   LYS A 135
        1054
               CA
 ATOM
                                                                               6
                                                               1.00 29.45
                                                       48.885
                                              17.971
               СВ
                   LYS A 135
                                     32.923
        1055
 ATOM
                                                               1.00 32.46
                                                                               6
                                                      48.638
                                     34.152
                                              17.131
                   LYS A 135
        1056
               CG
 ATOM
```

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MOTA	1057	CD	LYS A 135		33.965	15.734	49.221	1.00 29.67	6
			LYS A 135			15.703			
ATOM	1058	CE			34.234		50.716	1.00 30.09	6
ATOM	1059	NZ	LYS A 135		35.679	15.973	51.001	1.00 26.25	7
MOTA	1060	С	LYS A 135		33.513	19.516	46.993	1.00 30.22	6
ATOM	1061	0	LYS A 135		34.714	19.672	46.763	1.00 30.00	8
MOTA	1062	N	SER A 136		32.600	19.434	46.028	1.00 31.69	7
ATOM	1063	CA	SER A 136		32.995	19.489	44.619	1.00 32.88	6
ATOM	1064	CB	SER A 136		33.038	18.077	44.040	1.00 31.41	6
	1065	OG	SER A 136		33.882	17.241	44.810	1.00 35.41	
ATOM									8
ATOM	1066	С	SER A 136		32.097	20.347	43.727	1.00 33.55	6
MOTA	1067	0	SER A 136		31.921	20.031	42.553	1.00 36.11	8
ATOM	1068	N	ARG A 137		31.536	21.425	44.262	1.00 30.61	7
	1069	CA	ARG A 137		30.664	22.272	43.459	1.00 32.28	6
ATOM									
MOTA	1070	CB	ARG A 137		29.324	21.554	43.202	1.00 35.91	6
ATOM	1071	CG	ARG A 137		28.224	22.458	42.627	1.00 43.90	6
ATOM	1072	CD	ARG A 137		26.819	21.836	42.751	1.00 48.28	6
ATOM	1073	NE	ARG A 137	•	26.571	20.767	41.787	1.00 53.38	7
MOTA	1074	CZ	ARG A 137		26.150	20.960	40.538	1.00 55.30	6
ATOM	1075	NHl	ARG A 137		25.921	22.185	40.090	1.00 54.06	7
MOTA	1076	NH2	ARG A 137		25.969	19.922	39.728	1.00 58.96	7
ATOM	1077	С	ARG A 137		30.405	23.631	44.113	1.00 30.24	6
			ARG A 137		30.380	23.748			
MOTA	1078	0					45.338	1.00 23.11	8
MOTA	1079	N	ALA A 138		30.219	24.653	43.279.	1.00 27.33	7
ATOM	1080	CA	ALA A 138		29.944	26.000	43.757	1.00 27.36	6
ATOM	1081	CB	ALA A 138		30.149	26.997	42.645	1.00 27.57	6
	1082	č	ALA A 138		28.496	26.003	44.213	1.00 26.45	6
ATOM									
MOTA	1083	0	ALA A 138		27.747	25.083	43.865	1.00 27.30	8
ATOM	1084	N	ASN A 139		28.090	27.021	44.975	1.00 22.47	7
MOTA	1085	CA	ASN A 139		26.711	27.063	45.471	1.00 23.85	6
ATOM	1086	CB	ASN A 139		26.406	25.738	46.218	1.00 16.82	6
	1087	CG	ASN A 139		25.040	25.718			
MOTA							46.900	1.00 14.45	6
MOTA	1088		ASN A 139		24.019	26.084	46.319	1.00 13.39	8,
MOTA	1089	ND2	ASN A 139		25.018	25.249	48.139	1.00 20.08	7
MOTA	1090	С	ASN A 139		26.444	28.277	46.368	1.00 26.09	6
ATOM	1091	0	ASN A 139		27.239	28.600	47.260	1.00 27.50	8
								•	
MOTA	1092	N	GLY A 140		25.326	28.954	46.114	1.00 24.83	7
ATOM	1093	CA	GLY A 140		24.965	30.106	46.916	1.00 22.24	6
ATOM	1094	С	GLY A 140		25.991	31.211	46.890	1.00 22.35	6
MOTA	1095	0	GLY A 140		26.256	31.843	47.910	1.00 23.50	8
	1096	N	PHE A 141		26.570	31.437	45.717	1.00 25.60	7
ATOM									
MOTA	1097	CA	PHE A 141		27.582	32.476	45.518	1.00 26.47	6
MOTA	1098	CB	PHE A 141		27.204	33.765	46.258	1.00 28.05	6
ATOM	1099	CG	PHE A 141		25.925	34.391	45.792	1.00 28.61	6
ATOM	1100	CD1	PHE A 141		25.352	35.428	46.518	1.00 30.74	6
	1101		PHE A 141		25.312	33.975	44.620	1.00 29.10	6.
ATOM									
MC A	1102		PHE A 141		24.193	36.044	46.087	1.00 29.33	6
A'c'OM	1103	CE2	PHE A 141	4	24.150	34.583	44.177	1.00 31.03	6
A1CM	1104	CZ	PHE A 141		23.589	35.621	44.912	1.00 32.59	6
ATOM	1105	·C	PHE A 141		28.954	32.038	45.991	1.00 24.63	6
MOTA	1106	0	PHE A 141		29.938	32.727	45:733	1.00 29.72	8
ATOM	1107	N	CYS A 142		29.025	30.897	46.667	1.00 21.11	7
MOTA	1108	CA	CYS A 142		30.296	30.399	47.192	1.00 22.30	6
MOTA	1109	CB	CYS A 142		30.062	29.787	48.567	1.00 21.31	6
			CYS A 142		28.943	30.748	49.582	1.00 22.93	16
ATOM	1110	SG							
MOTA	1111	С	CYS A 142		31.017	29.366	46.326	1.00 22.13	6
ATOM	1112	0	CYS A 142		30.408	28.389	45.878	1.00 22.97	8
ATOM	1113	N	TYR A 143		32.317	29.573	46.111	1.00 23.09	7
	1114	CA	TYR A 143		33.129	28.632	45.335	1.00 23.05	6
ATOM									
MOTA	1115	CB	TYR A 143		34.063	29.365	44.375	1.00 21.60	6
MOTA	1116	CG	TYR A 143		33.377	30.379	43.487	1.00 24.09	6
ATOM	1117	CD1	TYR A 143		32.969	31.609	43.999	1.00 23.29	6
	1118		TYR A 143		32.365	32.555	43.199	1.00 23.26	6
MOTA							42.135		6.
MOTA	1119	CD2	TYR A 143		33.154	30.117		1.00 22.52	
ATOM	1120	CE2	TYR A 143		32.544	31.061	41.317	1.00 24.82	6
ATOM	1121	CZ	TYR A 143		32.153	32.281	41.857	1.00 27.55	6
ATOM	1122	ОН	TYR A 143		31.553	33.241	41.064	1.00 32.35	8
WT OLI									-

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		_			33.960	27.766	46.290	1.00 24.22	6
MOTA	1123		TYR A				45.998	1.00 24.58	8
ATOM	1124		TYR A		34.266	26.606			7
MOTA	1125	N	ILE A	144	34.327	28.329	47.437	1.00 23.83	
ATOM	1126	CA	ILE A	144	35.086	27.566	48.425	1.00 20.24	6
	1127		ILE A		36.547	27.982	48.453	1.00 17.27	6
ATOM					37.231	27.354	49.662	1.00 11.03	6
MOTA	1128		ILE A			27.603	47.110	1.00 14.93	6
MOTA	1129		ILE A		37.185			1.00 19.68	6
ATOM	1130	CD1	ILE A	144	38.601	28.028	46.946		
ATOM	1131	С	ILE A	144	34.495	27.703	49.815	1.00 21.77	6
ATOM	1132	0	ILE A	144	34.288	28.811	50.318	1.00 21.19	8
	1133		ASN A		34.212	26.555	50.424	1.00 23.00	7
MOTA			ASN A		33.616	26.508	51.750	1.00 20.92	6.
ATOM	1134				32.902	25.170	51.935	1.00 17.08	6
MOTA	1135		asn a				53.203	1.00 21.04	6
MOTA	1136		ASN A		32.079	25.125		1.00 20.97	8
ATOM	1137	OD1	ASN A	145	32.549	25.508	54.276		
ATOM	1138	ND2	ASN A	145	30.844	24.640	53.093	1.00 20.93	7
	1139		ASN A		34.706	26.669	52.806	1.00 19.68	б
MOTA			ASN A		35.201	25.679	53.351	1.00 20.64	. 8
MOTA	1140				35.079	27.911	53.100	1.00 16.28	7
MOTA	1141		ASN A			28.143	54.088	1.00 19.34	6
MOTA	1142	CA	ASN A		36.123				6
ATOM	1143	CB	ASN A	146	36.428	29.651	54.207	1.00 20.27	
ATOM	1144	CG	ASN A	146	35.292	30.444	54.795	1.00 18.05	6
	1145		ASN A		35.079	30.421	55.999	1.00 25.83	8
MOTA			ASN A		34.552	31.149	53.948	1.00 16.04	7
ATOM	1146				35.775	27.504	55.443	1.00 20.48	6
MOTA	1147	C	ASN A			27.027	56.151	1.00 19.88	8
ATOM	1148	0	ASN A		36.663		55.819	1.00 19.38	7
MOTA	1149	N	PRO A	147	34.482	27.485			6
ATOM	1150	CD	PRO A	147	33.312	28.068	55.135	1.00 17.48	
ATOM	1151	CA	PRO A	147	34.058	26.877	57.087	1.00 22.25	6
	1152	CB	PRO A		32.539	27.065	57.057	1.00 20.15	6
MOTA		CG	PRO · A		32.407	28.378	56.305	1.00 20.81	6
MOTA	1153		PRO A		34.443	25.383	57.188	1.00 26.89	6
MOTA	1154	C			35.066	24.954.		1.00 29.10	8
ATOM	1155	0	PRO A				56.176	1.00 25.88	7
ATOM	1156	N	ALA A		34.070	24.596			6
ATOM	1157	CA	ALA A	148	34.372	23.164	56.174	1.00 25.47	
ATOM	1158	CB	ALA A	148	33.670	22.468	55.009	1.00 21.84	6
	1159	С	ALA A	148	35.870	22.916	56.100	1.00 25.94	6
MOTA	1160	ŏ	ALA A		36.382	21.971	56.701	1.00 27.19	8
MOTA			VAL A		36.574	23.756	55.349	1.00 26.11	7
MOTA	1161	N			38.017	23.609	55.233	1.00 24.04	· 6
MOTA	1162	CA	VAL A			24.663	54.267	1.00 26.16	6
MOTA	1163	CB	VAL A		38.622			1.00 25.36	6
MOTA	1164	CG1	VAL A	149	40.135	24.476	54.158		6
ATOM	1165	CG2	VAL A	149	37.970	24.544	52.886	1.00 26.81	
	1166	С	VAL A		38.516	23.870	56.640	1.00 23.57	6
MOTA	1167	ō	VAL A		39.453	23.228	57.122	1.00 19.75	8
ATOM			GLY A		37.850	24.815	57.299	1.00 22.20	7
MOTA	1168	N			38.210	25.175	58.654	1.00 25.43	6
MOTĄ	1169	CA	GLY A	120			59.568	1.00 27.19	6
ATOM	1170	С	GLY A		38.130	23.975		1.00 27.05	8
ATOM	1171	0	GLY A	150	39.112	23.620	60.221		7
MOTA	1172	N	ILE A	151	36.959	23.348	59.618	1.00 25.56	
	1173	CA	ILE A		36.775	22.176	60.457	1.00 28.24	6
ATOM		CB	ILE. A		35.317	21.654	60.389	.1.00 29.41	6
ATOM	1174	CD	TIP A	151	35.251	20.215	60.869	1.00 26.02	6
MOTA	1175		ILE A	151		22.540	61.240	1.00 33.31	6
MOTA	1176	CG1	ILE A	151	34.394		CO 750	1.00 36.83	6
MOTA	1177	CD1	ILE A	151	34.255		60.759		6
MOTA	1178	С	ILE A	151	37.723	21.039	60.075	1.00 29.32	
	1179	ō	ILE A		38.340	20.420	60.947	1.00 29.08	8
MOTA		N	GLU A		37.843	20.769	58.778	1.00 29.91	7
MOTA	1180		GLU A		38.704		58.315	1.00 32.58	6
ATOM	1181	CA				_	56.802	1.00 35.07	6
ATOM	1182	CB	GLU A	T25	38.575			1.00 38.51	6
ATOM	1183	CG	GLU A	152	37.269		56.393		2
	1184	CD	GLU A	152	37.120		56.987	1.00 41.41	6
MOTA	1185	OE1		152	36.089	16.779	56.718	1.00 45.64	8
atom		_		152	38.030		57.723	1.00 40.09	8
ATOM	1186	OE2	OLU A	152	40.145		58.721	1.00 32.97	6
ATOM	1187	С	GLU A						8
ATOM	1188	0	GLU A	154	40.879	18.946			-
			•						

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Figure 17-19

ATCM	1189	N	TYR A 15:	3	40.541	21.170	58.765	1.00 33.90	7
ATOM	1190	CA			41.875		59.193	1.00 33.90	
MOTA	1191		•		42.019		59.058		6
ATOM	1192				43.280	-		1.00 34.88	6
ATOM	1193						59.639	1.00 38.03	6
	1194				44.498		58.948	1.00 42.34	6
ATOM					45.658			1.00 43.38	6
ATCM	1195		-		43.250		60.869	1.00 37.19	6
MCTA	1196				44.387	24.913	61.461	1.00 41.09	6
ATOM	1197	CZ	TYR A 153		45.587	24.860	60.704	1.00 43.34	6
ATOM	1198	OH	TYR A 153	1	46.696	25.480	61.241	1.00 44.86	8
ATOM	1199	С	TYR A 153		41.919	21.168	60.667	1.00 32.59	6
ATOM	1200	0	TYR A 153		42.867		61.120	1.00 32.39	
ATOM	1201		LEU A 154		40.869	21.556	61.397		8
ATOM	1202				40.730			1.00 30.10	7
MOTA	1203	CB				21.261	62.823	1.00 29.38	6
ATOM	1204	CG					63.378	1.00 28.60	6
					39.399	23.407	63.618	1.00 31.20	6
MOTA	1205		1 LEU A 154		37.991	23.833	64.041	1.00 28.53	6
ATOM	1206		2 LEU A 154		40.418	23.787	64.691	1.00 24.95	6
ATOM	1207	С	LEU A 154		40.732	19.772	63.146	1.00 29.56	6
MCTA	1208	0	LEU A 154		41.223	19.363	64.196	1.00 28,36	8
ATOM	1209	N	ARG A 155		40.174	18.958	62.256	1.00 31.95	7
MOTA	1210	CA	ARG A 155		40.134	17.522	62.499	1.00 33.00	6
ATOM	1211	CB	ARG A 155		39.127	16.847	61.561	1.00 33.13	6
ATOM	1212	CG	ARG A 155		37.708	17.368	61.769	1.00 32.84	6
ATOM	1213	CD	ARG A 155		36.678	16.719	60.863	1.00 32.84	
ATOM	1214	NE	ARG A 155	•	36.152	15.451			6
ATOM	1215	CZ	ARG A 155		35.195		61.363	1.00 33.98	7
ATOM	1216		L ARG A 155			14.760	60.741	1.00 37.93	6
	1217		2 ARG A 155		34.671	15.216	59.605	1.00 38.39	7
ATOM					34.732	13.631	61.259	1.00 38.67	7
ATOM	1218	C	ARG A 155		41.521	16.929	62.331	1.00 33.97	6
MOTA	1219	0	ARG A 155		41.869	15.941	62.985	1.00 32.95	8
ATOM	1220	N	LYS A 156		42.318	17.548	61.467	1.00 34.20	7
ATOM	1221	CA	LYS A 156		43.679	17.081	61.243	1.00 36.32	6
MOTA	1222	CB	LYS A 156		44.249	17.662	59.942	1.00 37.57	6
ATOM	1223	CG	LYS A 156		45.673	17.187	59.638	1.00 40.32	6
ATOM	1224	CD	LYS A 156		46.116	17.532	58.220	1.00 40.33	6
ATOM	1225	CE	LYS A 156		45.180	16.909	57.184	1.00 41.27	6
ATOM	1226	NZ	LYS A 156		45.015	15.435	57.364	1.00 37.92	7
MCTA	1227	С	LYS A 156		44.539	17.501	62.428	1.00 36.17	6
MOTA	1228	0	LYS A 156		45.582	16.905	62.699	1.00 34.53	. 8
MCTA	1229	N	LYS A 157		44.093	18.537	63.132	1.00 34.33	7
ATOM	1230	CA	LYS A 157		44.820	19.026	64.294		
MOTA	1231	СВ	LYS A 157		44.495	20.501		1.00 37.09	6
MCTA	1232	CG	LYS A 157				64.566	1.00 37.02	6
	1233	CD	LYS A 157		44.982	21.435	63.477	1.00 36.22	6
ATOM					46.468	21.231	63.239	1.00 37.91	6
MOTA	1234		LYS A 157		46.993	22.100	62.107	1.00 39.35	6
MCTA	1235	-JZ	LYS A 157		48.434	21.815	61.842	1.00 38.78	7
MCTA	1236	_	LYS A 157		44.498	18.178	65.515	1.00 35.61	6
ATOM	1237	0	LYS A 157		45.204	18.232	66.518	1.00 36.38	8
ATCM	1238	N	GLY A 158		43.433	17.392	65.431	1.00 34.37	7
MOTA	1239	CA	GLY A 158		43.097	16.537	66.552	1.00 38.08	6
ATOM	1240	С	GLY A 158		41.782	16.781	67.267	1.00 38.78	6
ATCM	1241	0	GLY A 158		41.460	16.053	68.208	1.00 41.07	8
ATOM	1242	N	PHE A 159		41.023	17.791	66.855	1.00 36.75	7
MOTA	1243	CA	PHE A 159		39.743	18.046	67.505		
ATOM	1244	CB	PHE A 159		39.246	19.459	67.213		6
MCTA	1245	CG	PHE A 159		40.115			1.00 32.65	6
	1246		PHE A 159			20.521	67.787	1.00 29.97	6
ATOM					41.404	20.724	67.297	1.00 30.20	6
MOTA	1247		PHE A 159		39.672	21.289	68.853	1.00 29.28	6
MCTA	1248		PHE A 159		42.241	21.680	67.862	1.00 28.96	6
MOTA	1249		PHE A 159		40.498	22.246	69.428	1.00 29.67	6
ATCM	1250	CZ	PHE A 159		41.785	22.442	68.931	1.00 30.59	6
ATCM	1251	С	PHE A 159		38.732	17.026	67.025	1.00 33.41	6
ATCM	1252	0	PHE A 159		38.664	16.716	65.838	1.00 31.61	8
ATCM	1253	N	LYS A 160		37.951	16.506	67.966	1.00 35.13	7
MOTA	1254	CA	LYS A 160		36.947	15.493	67.677	1.00 35.39	6
		-			J	20.200			J

ATOM	1255	CB	LYS A 16	0 7	7.342	14.198	68.389	1.00 36	: 43	6
	1256			-						
ATOM			· - - · · - ·	-	8.535	13.502	67.708	1.00 40		6
ATOM	1257	CD	LYS A 16	0 3	9.312	12.538	68.615	1.00 44	1.68	6
ATOM	1258	CE	LYS A 16	p 3	8.425	11.536	69.345	1.00 49	2 2 4	6
	1259									
MOTA					7.593	12.182	70.411	1.00 50		7
MOTA	1260	С	LYS A 16	3	5.524	15.927	68.027	1.00 35	94. ذ	6
MOTA	1261	0	LYS A 16	3	4.561	15.241	67.691	1.00 35	. 72	8
	1262	N	ARG A 16							
ATOM				_	5.399	17.058	68.718	1.00 34		7
ATOM	1263	CA	ARG A 16	1. 3	4.091	17.618	69.044	1.00 34	1.95	6
MOTA	1264	CB	ARG A 16	1 3	3.771	17.525	70.535	1.00 33	3 94	6
ATOM	1265	CG			3.427	16.132	70.992			
								1.00 38		6
MOTA	1266	CD			2.823	16.131	72.386	1.00 41	1.17	6
ATOM	1267	NE	ARG A 16	L 3	3.719	16.722	73.378	1.00 47	7 - 64	7
ATOM	1268	CZ	ARG A 16		4.912	16.233	73.705	1.00 47		6
MOTA	1269		1 ARG A 16		5.372	15.131	73.121	1.00 47		7
ATOM	1270	NH:	2 ARG [.] A 16	1 3	5.648	16.858	74.615	1.00 46	5.95	7
ATOM	1271	С	ARG A 16	. 3	4.113	19.076	68.598	1.00 34	1.58	6
ATOM	1272	0	ARG A 16		4.468	19.980	69.357	1.00 33		
										В
ATOM	1273	N	ILE A 16	د ؛	3.741	19.280	67.341	1.00 31	L.74	7
ATOM	1274	CA	ILE A 16	2 3	3.735	20.594	66.735	1.00 29	3.83	6
MOTA	1275	CB	ILE A 16		4.429	20.542	65.362	1.00 29		6
	•									
ATOM	1276	CG2			4.580	21.942	64.784	1.00 30		6
MOTA	1277	CG:	L ILE A 16	: 3	5.801	19.891	65.522	1.00 28	3.81	6
ATOM	1278	CD:	I ILE A 16	: 3	6.537	19.685	64.224	1.00 33	3 05	6
ATOM	1279	С	ILE A 16		2.300	21.050	66.560	1.00 29		
										6
ATOM	1280	0	ILE A 16		1.416	20.241	66.266	1.00 25	5.24	8
ATOM	1281	N	LEU A 16	3	2.081	22.351	66.745	1.00 30	0.00	7
ATOM	1282	CA	LEU A 16	. 3	0.754	22.945	66.617	1.00 30	1 48	6
	1283	CB	LEU A 16		0.236	23.406	67.992			
ATOM								1.00 32		6
MOTA	1284	CG	LEU A 16		8.934	24.229	68.044	1.00 3	1.21	6
ATOM	1285		L LEU A 16		7.804	23.494	67.326	1.00 33	1.58	6
ATOM	1286	CD2	LEU A 16	2	8.569	24.502	69.493	1.00 25	5 00	6
	1287	c	LEU A 16.		0.717	24.122				
ATOM							65.659	1.00 29		6
ATOM	1288	,O	LEU A 16	3:	1.596	24.980	65.654	1.00 29	€.72	8
ATOM	1289	N	TYR A 16	2:	9.675	24.157	64.846	1.00 29	9.68	7
ATOM	1290	CA	TYR A 16		9.500	25.244	63.899	1.00 2		6
	1291	CB								
MOTA			TYR A 16		9.512	24.688	62.470	1.00 2		6
ATOM	1292	CG	TYR A 16	2:	3.377	25.742	61.399	1.00 2	1.79	6
MOTA	1293	CD1	TYR A 164	3 (3.390	26.670	61.168	1.00 24	4.82	6
MOTA	1294	CE1			0.247	27.655	60.198	1.00 24		6
	1295	CD2								
ATOM					3.216	25.827	60.631	1.00 2		6
ATOM	1296	CE2	TYR A 164	28	3.065	26.808	59.662	1.00 2	5.67	5
MOTA	1297	CZ	TYR A 164	29	0.078	27.718	59.451	1.00 2	5.63	6
ATOM	1298	OH	TYR A 164		3.898	28.704	58.506	1.00 2		8
ATOM	1299	С	TYR A 164		3.149	25.907	64.218	1.00 28	3.38	6
ATOM	1300	Э	TYR A 164	27	7.119	25.225	64.277	1.00 29	€.43	8
ATOM	1301	N	ILE A 165	28	3.166	27.217	64.464	1.00 24	4 30	7
ATOM	1302	CA	ILE A 165		5.941	27.969	-64.754	1.00 2		6
MOTA	1303	CB	ILE A 165	2 t	.985	28.649	66.143	1.00 2	2.00	6
MOTA	1304	CG2	ILE A 165	25	.765	29.559	66.312	1.00 10	5.15·	6
MOTA	1305	CG1	ILE A 165	27	.033	27.567	67.240	1.00 20		6
	1306		ILE A 165				68.650	1 00 2	- 40	_
ATOM					1.185	28.101		1.00 1		6
ATOM	1307	С	ILE A 165	26	.784	29.010	63.657	1.00 24	1.45	6
MOTA	1308	o	ILE A 165	27	.605	29.921 [.]	63.506	1.00 23	3.17	8
ATOM	1309	N	ASP A 166		.709	28.871	62.895	1.00 24		7
ATOM	1310	CA	ASP A 166		.478	29.726	61.749	1.00 20		6
MOTA	1311	CB	ASP A 166		.314	28.809	60.548	1.00 1		6
ATOM	1312	CG	ASP A 166	25	.410	29.529	59.256	1.00 19	9.93	6
	1313		ASP A 166		.536	30.391	59.004	1.00 20		8
ATOM										0
ATOM	1314		ASP A 166		.366	29.231	58.491	1.00 1		8
MOTA	1315	C	ASP A 166		.290	30.670	61.895	1.00 2	2.79	б
ATOM	1316	0	ASP A 166	23	.134	30.256	61.826	1.00 22	2.70	8
-	1317	N	LEU A 167		.583	31.952	62.085	1.00 2		. 7
ATOM										
ATOM ·	1318	CA	LEU A 167		.536	32.954	62.250	1.00 2		6
MOTA	1319	CB	LEU A 167	23	.963	33.991	63.288	1.00 2	5.65	6
MOTA	1,32,0,	ÇG	LEU A 167	24	.364	33.463	64.674	1.00 2	5.75	6

	4227	an1			167		24.741	34.647	CE CE2	1.00 26.24	e
MOTA	1321		LEU						65.552		6
MOTA	1322	CD2	LEU	Α	167		23.225	32.661	65.302	1.00 23.45	6
MOTA	1323	С	LEU	Δ	167		23.162	33.660	60.951	1.00 26.37	6
							22.386	34.613	60.971	1.00 25.95	8
MOTA	1324	0	LEU								
MOTA	1325	N	ASP	Α	168		23.726	33.208	59.828	1.00 29.66	7.
ATOM-	1326	CA	ASP	Α	168		23.410	33.787	58.520	1.00 28.35	6
	1327		ASP				24.057	32.987	57.390	1.00 33.29	6
MOTA		CB									
MOTA	1328	CG	ASP	Α	168		23.937	33.676	56.037	1.00 35.38	6
ATOM ·	1329	OD1	ASP	Α.	168		24.892	34.388	55.659	1.00 39.48	8
	1330		-ASP				22.893	33.531	55.364	1.00 33.40	8
MOTA											
MOTA	1331	С	ASP				21.906	33.614	58.408	1.00 28.74	6
ATOM	1332	0	ASP	A.	168		21.354	32.648	58.948	1.00 26.21	8
MOTA	1333	N	ALA	A	169	-	21.239	34.524	57.711-	1.00 26.16	7
							19:793	34.415	57.579	1.00 24.39	6
MOTA	1334	CA	ALA								
MOTA	1335	CB	ALA	Α	169		19.233	35.640	56.879	1.00 22.75	6
MOTA	1336	С	ALA	Α	169		19.420	33.157	56.813	1.00 24.37	6 -
	1337	ō	ALA				18.266	32.752	56.824	1.00 22.34	8
ATOM											7
ATOM	1338	N	HIS				20.405	32.542	56.156	1.00 25.78	
MOTA	1339	CA	HIS	Α	170		20.180	31.327	55.375	1.00 25.20	6
ATOM	1340	CB	HIS	Α	170		20.667	31.501	53.936	1.00 25.76	6
			HIS				20.122	32.711	53.245	1.00 29.08	6
MOTA	1341	CG									
MOTA	1342	CD2	HIS	A	170		19.338	32.834	52.147	1.00 30.59	6
ATOM	1343	ND1	HIS	A	170		20.384	33.995	53.675	1.00 30.77	7
ATOM	1344		HIS				19.784	34.858	52.873	1.00 29.07	6
			HIS				19.143	34.180	51.939	1.00 32.19	7
ATOM	1345										
MOTA	1346	С	HIS	A	170		20.895	30.113	55.958	1.00 26.00	6
ATOM	1347	0	ĤΙS	A	170		21.913	30.234	56.637	1.00 25.76	8
ATOM	1348	N	HIS				20.349	28.939	55.658	1.00 27.29	7
									56.090	1.00 25.01	6
ATOM	1349	ÇA	HIS				20.893	27.655			
MOTA	1350	CB	HIS	A	171		19.934	26.532	55.663	1.00 24.93	6
MOTA	1351	CG	HIS	Α	171		20.468	25.148	55.889	1.00 26.56	6
ATOM	1352		HIS				20.674	24.123	55.028	1.00 22.34	6
									57.137	1.00 25.35	7
ATOM	1353		HIS				20.823	24.678			
MOTA	1354	CE1	HIS	Α	171		21.222	23.424	57.036	1.00 22.68	6
ATOM	1355	NE2	HIS	Α	171		21.140	23.062	55.767	1.00 24.13	7
	1356	С	HIS				22.267	27.413	55.471	1.00 24.74	6
MOTA								27.863	54.356	1.00 28.22	8
MOTA	1357	0	HIS				22.540				
MOTA	1358	N	CYS	А	172		23.131	26.705	56.190	1.00 23.03	7
ATOM	1359	CA	CYS	A	172		24.467	26.389	55.683	1.00 23.41	6
	1360	CB	CYS				25.497	26.474	56.812	1.00 19.31	6
MOTA										1.00 16.78	16
MOTA	1361 '	SG	CYS				25.005	25.631	58.318		
ATOM	1362	С	CYS	А	172		24.484	24.997	55.048	1.00 25.45	6
MOTA	1363	0 -	CYS	Α	172		25.203	24.098	55.483	1.00 24.47	8
	1364	N	ASP				23.664	24.839	54.015	1.00 26.67	7
MOTA								23.593		1.00 26.47	6
ATOM	1365	CA	ASP				23.542		53.269		
ATOM	1366	CB	ASP	Α	173		22.735	23.857	51.993	1.00 26.33	6
ATOM	1367	CG	ASP	Α	11.3		23.281	25.030	51.179	1.00 27.06	6
	1368		ASP			-	22.539	25.558	50.330	1.00 23.43	8
ATOM											8
MOTA	1369	OD2	ASP				24.454	25.417	51.372	1,.00 29.38	
MOTA	1370	С	ASP	Α	173		24.872	22.932	52.922	1.00 26.65	6
ATOM	1371	0	ASP	Α	173		24.940	21.708	52.784	1.00 28.38	8
		-								1.00 25.24	7
ATOM	1372	N	GLY				25.926	23.737	52.793		
ATOM	1373	CA	GLY	A	174		27.227	23.198	52.447	1.00 23.11	6
ATOM	1374	С	GLY	Α	174		27.896	22.505	53.612	1.00 25.64	6
	1375	ō	GLY				28.443	21.408	53.462	1.00 27.67	8
MOTA										1.00 24.29	7
ATCM	1376	N	VAL				27.848	23.144	54.778		
MOTA	1377	CA	VAL	À	175		28.459	22.602	55.989	1.00 22.20	6
ATOM	1378	CB	VAL	A	175		28.536	23.672	57.101	1.00 20.15	6
							29.449	23.192	58.218	1.00 20.11	6
atom	1379		VAL								
MOTA	1380	CG2	VAL	A	175		29.015	24.989	56.530	1.00 18.74	6
ATOM	1381	С	VAL	A	175		27.647	21.409	56.505	1.00 22.85	6
	1382	ō	VAL				28.173	20.512	57.173	1.00 20.07	8
MOTA									56.203	1.00 24.12	7
MOTA	1383	N	GLN				26.356	21.404			
ATOM ·	1384	ÇA	GLN				25.518	20.303	56.629	1.00 27.18	6
MOTA	1385	CB	GLN				24.045	20.611	56.355	1.00 32.86	6
		CG	GLN				23084	19.483	56.726	1.00 36.04	6
ATOM	1386	Ç.G	GTIM	^			22.009	17.403			-

	1207							
MOTA	1387	CI	GLN A 176	21.620	19.862	56.537	1.00 38.36	6
ATOM	1388	OF	E1 GLN A 176	21.113	20.782			
						57.185	1.00 38.59	8
ATOM	1389		2 GLN A 176	20.934	19.151	55.649	1.00 38.81	7
MOTA	1390	С	GLN A 176	25.956	19.083	55.841	1.00 27.70	6
MOTA	1391							
			GLN A 176		18.066	56.416	1.00 26.89	8
MOTA	1392	N	GLU A 177	25.951	19.194	54.519	1.00 27.96	7
MOTA	1393	CA	GLU A 177	26.343				
					18.062	53.698	1.00 31.16	6
ATOM	1394	CB	GLU A 177	26.395	18.460	52.220	1.00 30.37	6
MOTA	1395	CG	GLU A 177	26.353	17.256	51.287	1.00 36.20	
ATOM	1396							6
				26.273	17.626	49.818	1.00 40.70	6
ATOM	1397	OE	1 GLU A 177	27.322	17.967	49.234	1.00 46.78	8
ATOM	1398	OE	2 GLU A 177	25.155	17.590	49.250	1.00 39.39	
ATOM	1399	C	GLU A 177					8
				27.702	17.516	54.137	1.00 31.66	6
MOTA	1400	0	GLU A 177	27.868	16.317	54.356	1.00 32.81	8
ATOM	1401	N	ALA A 178	28.663	18.419	54.287	1.00 33.39	
ATOM	1402	CA						7
				30.026	18.072	54.673	1.00 31.63	6
ATOM	1403	CB	ALA A 178	30.830	19.338	54.856	1.00 30.96	6
ATOM	1404	С	ALA A 178	30.204	17.185	55.897	1.00 30.63	
								6
ATOM	1405	0	ALA A 178	31.032	16.276	55.876	1.00 27.95	8
ATOM	1406	N	PHE A 179	29.444	17.444	56.961	1.00 31.01	7
ATOM	1407	CA	PHE A 179	29.590	16.656			
						58.184	1.00 31.34	6
MOTA	1408	CB	PHE A 179	30.147	17.532	59.310	1.00 30.13	6
ATOM	1409	CG	PHE A 179	31.189	18.505	58.858	1.00 27.78	6
ATOM	1410	CD	1 PHE A 179	30.827	19.790			
						58.466	1.00 28.24	6
ATOM	1411	CD:		32.522	18.124	58.766	1.00 28.33	6
MOTA	1412	CE:	1 PHE A 179	31.778	20.688	57.988	1.00 26.68	6
ATOM	1413	CE:		33.487				
					19.013	58.285	1.00 28.79	6
ATOM	1414	CZ	PHE A 179	33.111	20.300	57.895	1.00 28.67	6
MOTA	1415	С	PHE A 179	28.300	16.003	58.664	1.00 32.06	6
ATOM	1416	0	PHE A 179	28.218	15.542			
						59.803	1.00 30.58	8
MOTA	1417	N	TYR A 180	27.305	15.960	57.787	1.00 34.25	7
ATOM	1418	CA	TYR A 180	26.001	15.377	58.099	1.00 38.60	6
ATOM	1419	CB	TYR A 180	25.062	15.605	56.911		
							1.00 38.99	6
ATOM	1420	CG	TYR A 180 -		15 <i>:</i> 453	57.220	1.00 37.91	6
ATOM .	1421	CD1	L TYR A 180	22.938	14.232	57.064	1.00 35.83	6
ATOM	1422	CE1	TYR A 180	21.589	14.103	57.373		
	1423						1.00 39.20	6
ATOM		CD2		22.861	16.543	57.694	1.00 37.56	6
MOTA	1424	CE2	TYR A 180	21.518	16.430	58.007	1.00 40.28	6
MOTA	1425	CZ	TYR A 180	20.882	15.211	57.848	1.00 41.92	
	1426	OH						6
ATOM			TYR A 180	19.549	15.110	58.188	1.00 43.41	8
ATOM	1427	C	TYR A 180	26.133	13.884	58.382	1.00 40.28	6
ATOM	1428	0	TYR A 180	25.158	13.192	58.680	1.00 39.27	8
MOTA	1429	N	ASP A 181			_		
				27.363	13.402	58.319	1.00 43.51	7
ATOM	1430	CA	ASP A 181	27.638	11.994	58.519	1.00 45.89	6
ATOM	1431	CB	ASP A 181	28.414	11.487	57.303	1.00 51.00	6
	1432	CG	ASP A 181					
ATOM				28.830	10.050	57.436	1.00 56.84	6
MOTA	1433	OD1	ASP A 181	29.637	9.750	58.345	1.00 59.47	8
ATOM	1434	OD2	ASP A 181	28.348	9.221	56.629	1.00 60.73	8
MOTA	1435	С	ASP A 181					
				28.398	11.665	59.804	1.00 44.75	6
ATOM	1436	0	ASP A 181	28.257	10.568	60.350	1.00 44.69	8
MOTA	1437	N	THR A 182	29.194	12.506	60.298	1.00 41.26	7
	1438	CA						
ATOM			THR A 182	29.975	12.337	61.495	1.00 39.51	6
ATOM	1439	CB	THR A 182	31.408	12.881	61.355	1.00 39.19	6
ATOM	1440	0G1	THR A 182	32.171	12.508	62.505	1.00 37.82	8
	1441		THR A 182					
ATOM				31395	14.397	61.232	1.00 40.12	6
MOTA	1442	С	THR A 182	29.370	12.910	62.759	1.00 38.58	6
ATOM	1443	0	THR A 182	28.609	13.876	62.716	1.00 41.24	8
	1444	N	ASP A 183					-
ATOM				29.712	12.304	63.890	1.00 37.39	7
ATOM	1445	CA	ASP A 183	29.211	12.773	65.171	1.00 39.24	6
ATOM	1446	CB	ASP A 183	28.824	11.588	66.061	1.00 40.31	6
	1447	CG	ASP A 183					
ATOM				30.010	10.723	66.433	1.00 41.64	6
ATOM	1448	ODI	ASP A 183	30.725	10.268	65.520	1.00 42.53	8
ATOM	1449	OD2	ASP A 183	30.221	10.494	67.640	1.00 42.46	8
ATOM	1450	c	ASP A 183	30.286			-	
					13.621	65.853	1.00 40.34	6
ATOM	1451	0	ASP A 183	30.109	14.071	66.983	1.00 42.07	8
ATOM	1452	N	GLN A 184	31.400	13.830	65.154	1.00 39.29	7
<u></u>								•

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MOTA	1453	C2	A GLN A 184	32.506	14.635	65.671	1.00 37.08	6
ATOM	1454	CI	3 GLN A 184	33.830				6
MOTA	1455	i co	GLN A 184	34.229				6
MOTA	145€	CI	GLN A 184	35.599				6
MOTA	1457	OF	E1 GLN A 184	35.853	12.704			8
MOTA	1458	NE.	E2 GLN A 184	36.490				7
MOTA	1459	C	GLN A 184	32.222				6
MOTA	1460	0	GLN A 184	32.803		66.034		8
MOTA	1461	. N	VAL A 185	31.329		64.456		7
MOTA	1462	CA		30.984				6
MOTA	1463			31.308		62.641	1.00 33.03	6
MOTA	1464		1 VAL A 185	31.009				6
ATOM	1465		2 VAL A 185	32.773	17.738	62.357		6
MOTA	1466		VAL A 185	29.508	17.972	64.360	1.00 31.51	6
MOTA	1467		VAL A 185	28.680	17.124	64.038		8
ATOM	1468		PHE A 186	29.185		64.946		7
MOTA	1469			27.798	19.463	65.207	1.00 31.44	6
MOTA	1470	CB		27.524	19.532	66.716	1.00 30.96	6
MOTA	1471	CG		26.059	19.617	67.066	1.00 31.59	6
MOTA	1472		1 PHE A 186	25.552	18.901	68.153	1.00 30.54	6
ATOM	1473	CD		25.179	20.395	66.308	1.00 31.50	6
MOTA	1474		1 PHE A 186	24.191	18.951	68.478	1.00 33.28	6
MOTA	1475		2 PHE A 186	23.815	20.457	66.622	1.00 33.04	6
MOTA	1476	cz	PHE A 186	23.318	19.733	67.708	1.00 32.35	6
ATOM	1477	C	PHE A 186	27.490	20.798	64.551	1.00 30.37	6
MOTA MOTA	1478 1479	O N	PHE A 186 VAL A 187	28.189	21.789	64.751	1.00 31.32	8
ATOM	1480	CA	VAL A 187	26.435	20.809	63.752	1.00 31.14	7
MOTA	1481	CB	VAL A 187	26.024 26.018	22.015	63.063	1.00 32.05	6
ATOM	1482		1 VAL A 187	25.574	21.805 23.081	61.525	1.00 33.54	6
ATOM	1483		2 VAL A 187	27.420	21.389	60.813 61.056	1.00 32.07	Ę
ATOM	1484	C	VAL A 187	24.638	22.439	63.524	1.00 35.44	6
ATOM	1485	0	VAL A 187	23.666	21.686.		1.00 31.47 1.00 29.06	6 8
ATOM	1486	N	LEU A 188	24.579	23.638	64.090	1.00 29.44	7
MOTA	1487	CA	LEU A 188	23.336	24.228	64.551	1.00 29.39	6
MOTA	1488	CB	LEU A 188	23.433	24.665	66.009	1.00 29.62	6
MOTA	1489	CG	LEU A 188	22.293	25.589	66.458	1.00 27.92	6
MOTA	1490	CD1	LEU A 188	20.970	24.844	66.414	1.00 25.87	6
ATOM	1491		LEU A 188	22.574	26.107	67.861	1.00 27.69	6
ATOM	1492	C	LEU A 188	23.161	25.454	63.675	1.00 31.89	6
MOTA	1493	0	LEU A 188	24.130	26.175	63.388	1.00 31.50	8
ATOM	1494	N	SER A 189	21.929	25.700	63.250	1.00 29.93	7
MOTA	1495 1496	CA	SER A 189	21.682	26.831	62.390	1.00 24.65	6
ATOM ATOM	1497	CB OG	SER A 189 SER A 189	21.873	26.411	60.942	1.00 22.40	6
ATOM	1498	C	SER A 189	21.585	27.485	60.083	1.00 19.12	8
ATOM	1499	ŏ	SER A 189	20.716 19. <i>2</i> 96	27.462	62.540	1.00 27.00	6
ATOM	1500	N	LEU A 190	20.521	26.774 28.783	62.577	1.00 26.72	8
ATOM	1501	CA	LEU A 190	19.096	29.554	62.669 62.735	1.00 27.41	7
ATOM	1502	CB	LEU A 190	19.185	30.682	63.771	1.00 29.68	6
MOTA	1503	CG	LEU A 190	19.108	30.366		1.00 29.84 1.00 26.79	6 6
ATOM	1504		LEU A 190	19.020	31.662	66.045	1.00 28.79	6
ATOM	1505		LEU A 190	17.881	29.549	65.546	1.00 27.63	6
ATOM	1506	С	LEU A 190	19.046	30.141	61.329	1.00 29.58	6
ATOM	1507	0	LEU A 190	20.084	30.525	60.790	1.00 32.40	8
ATOM	1508	N	HIS A 191	17.864	30.206	60.727	1.00 29.61	7
MOTA	1509	CA	HIS A 191	17.766	30.726	59.368	1.00 29.72	6
MOTA	1510	CB	HIS A 191	18.595	29.839	58.432	1.00 26.47	6
MOTA	1511	CG	HIS A 191	18.225	28.392	58.504	1.00 28.18	6
MOTA	1512		HIS A 191	18.918	27.313	58.940	1.00 28.88	6
MOTA	1513		HIS A 191	16.989	27.921	58.118	1.00 31.00	7
ATCM	1514		HIS A 191	16.938	26.614	58.312	1.00 30.54	6
ATOM	1515		HIS A 191	18.095	26.220	58.810	1.00 27.21	7
ATOM	1516	C	HIS A 191	16.329	30.812	58.856	1.00 28.05	6
ATOM	1517	0	HIS A 191		30.411	59.535	1.00 27.81	8
MOTA	1518	N	GLN A 192	16.183	31.346	57.649	1.00 29.39	7

OTA	M 1519 CA GLN A 192	14 00	c 21 404	F		
ATC		14.88				6
		15.01		55.796	1.00 24.94	ϵ
ATC		15.62	2 33.773	56.124	1.00 21.66	6
ATC	M 1522 CD GLN A 192	15.70				
ATO		14.684				6
ATO						8
		16.91		54.434	1.00 22.97	7
ATO		14.43	30.104	56.570		6
ATO	M 1526 O GLN A 192	15.157				
ATO		13.249				8
ATO						7
		12.751			1.00 33.28	6
ATO		- 11.264	28.249	56.961		6
ATO	M 1530 OG SER A 193	10.786		56.540		
ATO		12.974		50.540		8
				55.171	1.00 34.79	6
ATO		12.775		54.356	1.00 33.74	8
ATO		13.404	26.938	54.803	1.00 37.57	7
ATO:	M 1534 CD PRO A 194	13.689		55.658	1.00 38.89	
ATO						6
		13.654		53.403	1.00 37.81	6
ATO:		14.248	25,194	53.498	1.00 39.30	6
ATO	M 1537 CG PRO A 194	14.840	25.163	54.916	1.00 39.09	6
ATO	M 1538 C PRO A 194	12.340		52.638		
ATO					1.00 36.81	6
		12.317		51.425	1.00 34.09	8
OTA		11.246	26.835	53.364	1.00 39.25	7
ATO	M 1541 CA GLU A 195	9.928	26.866	52.750	1.00 41.54	6
ATO	1 1542 CB GLU A 195	8.843		53.812		
ATO					1.00 45.84	6
		8.360		54.608	1.00 53.30	6
ATON		7.160	28.502	53.960	1.00 55.91	6
OTA	1 1545 OE1 GLU A 195	6.735	29.571	54.461	.1.00 55.14	8
OTA		6.631		52.956		
ATON					1.00 57.75	8
		9.700		52.047	1.00 39.50	6
ATON		8.651	28.431	51.452	1.00 40.21	8
ATOM	1 1549 N TYR A 196	10.689	29.096	52.098	1.00 37.13	7.
ATOM	1 1550 CA TYR A 196	10.549	30.379	51.434	1.00 35.50	6
ATOM						
		9.602	31.274	52.245	1.00 36.36	6
ATOM		10.175	31.816	53.538	1.00 37.28	6
ATOM		11.120	32.848	53.527	1.00 35.42	6
ATOM	1554 CE1 TYR A 196	11.637	33.366	54.706	1.00 33.10	
ATOM		9.764				6
			31.311	54.776	1.00 36.75	6
ATOM		10.279	31.825	55.968	1.00 35.26	6
ATOM	1557 CZ TYR A 196	11.213	32.856	55.922	1.00 35.84	6
ATOM	1558 OH TYR A 196	11.704	33.401	57.087.	1.00 37.09	8
ATOM		11.878				
	· · · · · · · · · · · · · · · · · · ·		31.097	51.188	1.00 34.89	6
ATOM		11.896	32.256	50.764	1.00 31.61	8
ATOM		12.991	30.416	51.437	1.00 34.39	7
ATOM	1562 CA ALA A 197	14.297	31.041	51.242	1.00 34.82	6
ATOM		14.684				
			31.826	52.489	1.00 32.48	6
MOTA		15.418	30.075	50.887	1.00 36.59	6
ATOM		15.407	28.903	51.291	1.00 37.46	8
ATOM	1566 N PHE A 198	16.388	30.584	50.133	1.00 36.22	7
ATOM	1567 CA PHE A 198	17.548	29.802	49.722	1.00 37.68	
ATOM	1568 CB PHE A 198					6
		18.597	30.729	49.109	1.00 40.89	6
MOTA	1569 CG PHE A 198	19.810	30.013	48.578	1.00 43.59	6
ATOM	1570 CD1 PHE A 198	19.783	29.404	47.325	1.00 44.74	6
ATOM	1571 CD2 PHE A 198	20.970	29.929	49.336		
ATOM	1572 CE1 PHE A 198				1.00 41.86	6
	1572 CEI PRE A 198	20.894	28.729·	46.833	1.00 41.42	6
ATOM	1573 CE2 PHE A 198	22.079	29 251	48.849	1.00 43.30	6
MOTA	1574 CZ PHE A 198	22.040	28.652	47.595	1.00 41.86	6
ATOM	1575 C PHE A 198	18.139		50.967		
	1576 O PHE A 198		29.140		1.00 37.00	6
ATOM		18.166	29.754	52.036	1.00 36.43	8
MOTA	1577 N PRO A 199	18.641	27.892	50.848	1.00 37.63	7
ATOM	1578 CD PRO A 199	19.298	27.238	51.997	1.00 35.29	6
ATOM	1579 CA PRO A 199	18.727		49.673		
					1.00 36.52	6
ATOM		19.702		50.138	1.00 34.96	6
ATOM	1581 CG PRO A 199	19.281	25.770	51.565	1.00 34.57	б
ATOM	. 1582 C PRO A 199	17.409		49.222	1.00 35.72	6
MOTA	1583 O PRO A 199	17.386		48.225	1.00 37.36	
	- ·				1.00 37.36	8
ATOM		16.331	26.638	49.962	1.00 33.78	7
			•	-	•	

ATOM	1585	CA	PHE A	200		15.00	4 :	26.090	10	662	1 0	0 32.1	-
ATOM			PHE A			14.56		26.381					
ATOM			PHE A			14.60				222		0 28.3	_
ATOM			1 PHE A				_	27.827		835		26.2	
						L5.74		8.385	-	296		0 24.8	
ATOM			2 PHE A			13.46		8.623		966	1.0	28.0	4 6
ATOM			L PHE A			5.76		9.712		882	1.00	25.6	
MOTA			PHE A		1	.3.47	52	9.955	47.	557	1.00	27.0	
MOTA			PHE A		1	4.62	6 3	0.498		013		24.9	
MOTA	1593	С	PHE A	200	1	4.94	72	4.574		842		32.6	
ATOM	1594	0	PHE A	200		3.92		4.033		264		31.2	-
MOTA	1595	N	GLU A			6.043		3.896		499	1.00	31.2	2 8
ATOM	1596		GLU A			6.128						33.6	
ATOM	1597	СВ	GLU A			7.213	_	2.438	49.			30.8	
	1598							1.931		637-		32.9	-
ATOM		CG	GLU A			6.879		2.182	47.			33.5	
ATOM	1599	CD	GLU A			8.012		1.864	46.3	232	1.00	34.5	6 6
MOTA	1600		GLU A		1	8.396	5 2	0.678	46.3	117	1.00	36.3	5 8
ATOM	1601	OE2	GLU A	201	1	8.523	2	2.814	45.0	605		36.5	
MOTA	1602	С	GLU A .	201	1	6.369	2	1.911	50.9			28.5	
ATOM	1603	0	GLU A	201		5.537		1.199	51.5			28.9	_
MOTA	1604	N	LYS A			7.511		2.239	51.9				-
MOTA	1605	CA	LYS A			7.795						31.6	
MOTA	. 1606	СВ	LYS A					1.780	52.9			32.3	-
ATOM	1607	CG	LYS A			9.276		1.432	53.0			36.9	-
						9.789		0.226	52.3			43.7	
MOTA	1608	CD	LYS A			0.212		0.590	50.8	391	1.00	49.3	16
MOTA	1609	CE	LYS A 2			0.952		9.428	50.2	227	1.00	49.34	
MOTA	1610	NZ	LYS A 2		2:	1.504	19	9.802	48.8	395	1.00	49.89	
ATOM	1611	C	LYS A 2		1'	7.421	22	2.849	53.9	37	1.00	30.5	
MOTA	1612	0	LYS A 2	202	. 16	5.877	23	3.890	53.5			26.5	
MOTA	1613	N	GLY A 2	203		7.710		2.571	55.2			30.59	
MOTA	1614	CA	GLY A 2	203		7.422		3.519	56.2			30.24	
ATOM	1615	С	GLY A 2			5.216		3.210	57.1			29.23	
ATOM	1616	Ō	GLY A 2			5.915		3.975					
ATOM	1617	N	PHE A 2			5.526			58.0			32.90	-
ATOM	1618	CA	PHE A 2					2.104	56.8			26.32	•
	1619					1.344		779	57.6			23.25	
ATOM		CB	PHE A 2			.366		.917	56.8		1.00	21.25	6
ATOM	1620	CG	PHE A 2			.855	21	.573	55.6	35	1.00	18.60	6
MOTA	1621		PHE A 2		13	.605	21	.560	54.4	61	1.00	16.43	6
MOTA	1622		PHE A 2		11	.654	22	.273	55.6	64	1.00	14.82	
ATOM	1623		PHE A 2		13	.168	22	.245	53.3	33	1.00	16.91	
MOTA	1624	CE2	PHE A 2	04	11	.206		.962	54.5			15.28	
MOTA	1625	CZ	PHE A 2	04	11	.965		.952	53.3			18.34	
ATOM	1626	С	PHE A 2	04		.626		.094	58.9			23.72	
ATOM	1627		PHE A 2			.578		.318	59.1			22.68	_
MOTA	1628		LEU A 2			.760		.376	59.9				
ATOM	1629		LEU A 2			.877						20.94	•
ATOM	1630		LEU A 20					.818	61.2			24.83	-
	1631					.678		.259	52.1			21.29	-
MOTA			LEU A 20			.672			33.50		1.00	22.67	6
MOTA	1632		LEU A 20			.011	21	.182	74.24	45	1.00	19.76	6
ATOM			LEU A 20		11	.478	21	. 456	64.27	75	1.00	20.62	6
MOTA			LEU A 20		14	.002	19	. 293	61.30	3	1.00	28.79	
ATOM			LEU A 20		14	.443	18	.730	62.31	LO		28.59	. 8
MOTA	1636	N C	GLU A 20)6	13	. 625		. 628	60.21		1.00		7
ATOM	1637	CA C	GLU A 20	16		. 693		.166	60.14		1.00		6
ATOM			LU A 20			.736		.616	59.07		1.00		
ATOM			LU A 20			.284							6
ATOM			LU A 20			014		.060	59.20		1.00		6
								. 390	58.51		1.00		6
ATOM .			LU A 20			972		.027	58.79		1.00		8
ATOM			LU A 20			839		.786	57.66		1.00		8
ATOM			LU A 20		15.	114	16.	674	59.84		1.00	40.00	6
MOTA	1644 (0 G	LU A 20	6	15.	483		541	60.18		1.00		8
MOTA	1645 1	N G	LU A 20	7		903		536	59.21		1.00		7
MOTA			LU A 20			286		219	58.87		1.00		6
MOTA	_		LU A 20			776		242					
ATOM .		_	LU A 20						57.85		1.00		6
						983		158	56.55		1.00		6
MOTA			LU A 20			978			55.77		1.00		6
MOTA	1650 C	CET C	LU A 20'			071	20.	016	55.53	7	1.00	55.44	8
		•			•				-				

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MOTA	1651	OE	2 GLU A 207	15.870	19.891	55.389	1.00 34.62	8
MOTA	1652	C	GLU A 207	18.139	17.239	60.134	1.00 36.18	6
ATOM	1653	0	GLU A 207	18.560	18.303	60.590	1.00 34.81	8
ATOM	1654	N	ILE A 208	18.381	16.059	60.701	1.00 34.45	7
MOTA	1655	CA	ILE A 208	19.164	15.965	61.932	1.00 38.53	6
MOTA	1656	CB	ILE A 208	18.260	15.511	63.117	1.00 41.12	6
MOTA	1657	CG		19.097	15.273	64.375	1.00 41.36	6
MOTA	1658	CG		17.193	16.581	63.383	1.00 42.21	6
MOTA	1659	CD		16.291	16.286	64.560	1.00 44.81	6
MOTA	1660	C	ILE A 208	20.407	15.074	61.876	1.00 36.66	6
MOTA	1661	0	ILE A 208	21.243	15.110	62.775	1.00 34.03	8
ATOM	1662	N	GLY A 209	20.540	14.284	60.822	1.00 36.80	7
MOTA	1663	CA	GLY A 209	21.703	13.428	60.728	1.00 38.99	6
MOTA	1664	C	GLY A 209	21.509	12.246	59.805	1.00 40.93	6
ATOM	1665	0	GLY A 209	20.477	12.124	59.145	1.00 40.26	8
ATOM	1666	N	GLU A 210	22.508	11.370	59.775	1.00 42.16	7
MOTA	1667	CA	GLU A 210	22.492	10.185	58.930	1.00 43.30	6
ATOM	1668	CB	GLU A 210	22.810	10.586	57.488	1.00 47.08	6
ATOM	1669	CG	GLU A 210	22.826	9.453	56.478	1.00 53.90	6
MOTA	1670	CD	GLU A 210	23.256	9.915	55.089	1.00 56.27	6
ATOM	1671 1672	OE:		24.412	10.371	54.941	1.00 56.19	8
MOTA	1673	C	GLU A 210	22.437 23.583	9.826	54.145	1.00 60.28	8
ATOM ATOM	1674	ō	GLU A 210	24.750	9.276 9.457	59.473 59.152	1.00 41.47	6 8
ATOM	1675	N	GLY A 211	23.203	8.307	60.299	1.00 43.97	7
ATOM	1676	CA	GLY A 211	24.181	7.405	60.885	1.00 37.34	6
ATOM	1677	C	GLY A 211	24.642	7.952	62.224	1.00 37.84	6
ATOM	1678	O	GLY A 211	23.820	8.408	63.019	1.00 37.30	8
ATOM	1679	N	LYS A 212	25.948		62.485	1.00 38.52	7
ATOM	1680	CA	LYS A 212	26.490	8.440	63.733	1.00 38.29	6
ATOM	1681	CB	LYS A 212	28.020	8.359	63.731	1.00 40.54	6
ATOM	1682	CG	LYS A 212	28.570	6.950	63.675	1.00 46.39	6
ATOM	1683	CD	LYS A 212	28.149	6.147	64.910	1.00 51.59	6
MOTA	1684	CE	LYS A 212	28.556	4.676	64.809	1.00 52.77	б
ATOM	1685	NZ	LYS A 212	30.030	4.478	64.662	1.00 55.48	7
MOTA	1686	C	LYS A 212	26.061	9.897	63.866	1.00 37.68	6
MOTA	1687	0	LYS A 212	25.814	10.389	64.962	1.00 34.75	8
ATOM	1688	N	GLY A 213	25.956	10.574	62.728	1.00 38.89	7
ATOM	1689 1690	CA C	GLY A 213 GLY A 213	25.577	11.975	62.724	1.00 43.58	6
ATOM	1691	o	GLY A 213	24.126 23.737	12.295 13.464	63.020 63.024	1.00 43.99 1.00 44.67	6 8
ATOM ATOM	1692	N	LYS A 214	23.321	11.268	63.265	1.00 46.02	7
ATOM	1693	CA	LYS A 214	21.907	11.467	63.562	1.00 45.61	6
ATOM	1694	CB	LYS A 214	21.168	10.130	63.469	1.00 47.77	6
ATOM	1695	CG	LYS A 214	19.675	10.252	63.249	1.00 49.25	6
ATOM	1696	CD	LYS A 214	19.078	8.901	62.911	1.00 51.64	6
ATOM ·	1697	CE	LYS A 214	17.637	9.038	62.440	1.00 54.30	6
MOTA	1698	NZ	LYS A 214	17.030	7.727	62.034	1.00 56.09	7
ATOM	1699	С	LYS A 214	21.809	12.047	64.970	1.00 44.22	6
ATOM	1700	0	LYS A 214	22.210	11.410	65.942	1.00 45.04	8
ATOM	1701	N	GLY A 215	21.292	13.266	65.074	1.00 42.89	7
MOTA	1702	CA	GLY A 215	21.193	13.904	66.373	1.00 40.20	6
MOTA	1703	С	GLY A 215	22.295	14.931	66.600	1.00 39.90	6
MOTA	1704	0	GLY A 215	22.356	15.548	67.668	1.00 40.12	8
MOTA	1705	N	TYR A 216	23.175	15.111	65.612	1.00 38.49	7
ATOM	1706	CA	TYR A 216	24.261	16.086	65.726	1.00 35.91	6
ATOM	1707	CB	TYR A 216	25.632	15.421	65.618	1.00 36.47	6
MOTA	1708	CG	TYR A 216	25.935	14.461	66.738	1.00 39.54	6
MOTA	1709		TYR A 216	25.296	13.220	66.810	1.00 40.62	6
ATOM	1710		TYR A 216	25.561	12.333	67.849	1.00 41.58	6
ATOM	1711		TYR A 216	26.852	14.795	67.739	1.00 39.20	6
MOTA	1712		TYR A 216	27.124	13.915	68.786	1.00 40.87	6 6
ATOM	1713	CZ OH	TYR A 216 TYR A 216	26.475 26.743	12.685	68.833	1.00 41.86 1.00 43.04	8
ATOM	1714 1715	C	TYR A 216	24.182	11.802 17.215	69.852 64.709	1.00 43.04	6
ATOM ATOM	1716	ō	TYR A 216	25.194	17.832	64.375	1.00 34.21	8
MIUM	1,10	9	8 410	ムン・エフセ	11.032	Ja.3/3	1.00 33,37	-

MOTA	171	7 N	ASN	7 A 217	22.	976	17.471	64.212	1.00	33.83	7
MOTA	171	8 C.	A ASN	I A 217		726		63.267		30.20	
ATOM	171			I A 217							6
						699		61.823		27.74	6
MOTA	172			I A 217		457		60.826	1.00	25.61	6
ATOM	172	1 0	D1 ASN	I A 217	21.	354	19.705	60.719	1.00	25.00	8
MOTA	172	2 N	D2 ASN	A 217	23.	501		60.103		30.43	7
ATOM	172			A 217		369					
								63.645		29.09	6
MOTA	172			A 217	20.	433	18.351	63.885	1.00	26.93	8
MOTA	172	5 N	LEU	A 218	21.	263	20.440	63.710	1.00	27.19	7
MOTA	172	S C	A LEU	A 218	·20.	010	21.071	64.089		25.33	6
ATOM	172			A 218							
						026	21.379	65.590		23.23	б
MOTA	1728		_	A 218	18.	729	21.704	66.346	1.00	21.00	6
MOTA	1729	CI	Ol LEU	A 218	19.	100	22.313	67.695	1.00	18.62	6
ATOM	1730	CI	2 LEU	A 218		872		65.583		18.48	
ATOM	1731			A 218							6
					19.		22.368	63.325		25.04	6
MOTA	1732			A 218	20.	596	23.287	63.415	1.00	25.23	8
ATOM	1733	N	asn	A 219	18.	681	22.436	62.584	1.00	28.44	7
ATOM	1734	CA	ASN	A 219	18.	310	23.636	61.829		28.76	6
ATOM	1735			A 219	17.						
							23.298	60.417		25.69	6
ATOM	1736			A 219	18.		22.408	59.646	1.00	26.10	6
MOTA	1737	OD	1 ASN	A 219	19.	927	22.708	59.505	1.00	28.53	8
ATOM	1738	ND	2 ASN	A 219	18.3	220	21.311	59.114		26.97	7
ATOM	1739			A 219	17.		24.248	62.582		-	
	1740			A 219						31.96	6
MOTA					16.		23.539	63.246		34.84	8
ATOM	1741			A 220	16.9	952	25.556	62.472	1.00	32.96	7
MOTA	1742	CA	ILE	A 220	15.8	326	26.196	63.129	1.00	32.50	6
MOTA	1743	CB	ILE	A 220	16.2		27.037	64.350		32.32	6
ATOM	1744	CG		A 220	15.0						
	1745						27.644	65.014		29.46	6
MOTA		CG.		A 220	16.9		26.160	65.374	1.00	29.65	6
MOTA	1746	CD.	1 ILE	A 220	16.0	080	25.138	66.027	1.00	28.65	6
MOTA	1747	С	ILE	A 220	15.1	140	27.106	62.123	1.00	35.36	6
ATOM	1748	0	TLE	A 220	15.4		28.290	62.009		35.52	8
ATOM	1749	N		A 221							
					14.1		26.553	61.359		36.87	7
MOTA	1750	CD		A 221 .	13.7		25.158	61.359		35.12	6
ATOM	1751	CA	PRO	A 221	13.4	45	27.318	60.356	1.00	35.41	6
ATOM	1752	CB	PRO	A 221	12.5	09	26.262	59.767		35.68	6
ATOM	1753	CG		A 221	13.3		24.992	59.911			
										33.86	6
MOTA	1754	C		A 221	12.6		28.437	61.053	1.00	34.37	6
ATOM	1755	0	PRO .	A 221	12.0	14	28.199	62.043	1.00	38.79	8
ATOM	1756	N	LEU .	A 222	12.8	15	29.655	60.547	1.00	34.76	7
ATOM	1757	CA	LEU 2	A 222	12.1		30.796	61.166		33.87	6
ATOM	1758	CB		A 222							
					13.1		31.735	61.798		35.13	6
MOTA	1759	CG		A 222	14.1		31.163	62.876	1.00	33.07	6
MOTA	1760		LEU 2		15.2	34	32.150	63.154	1.00	34.04	6
MOTA	1761	CD2	LEU A	A 222	13.3	12	30.856	64.141		32.39	6
ATOM	1762	С		A 222	11.2		31.567	60.157	1.00	22.05	6
	1763	ō		222							
MOTA					11.6		31.740	59.000	1. 0		· 8
ATOM	1764	N	PRO A		10.1	27	32.060	60.601	1.00	30.97	7
ATOM	1765	CD	PRO A	A 223	9.6	06	31.913	61.972	1.00	32.34	6
ATOM	1766	CA	PRO A	223	9.1	73	32.818	59.789	1.00	30 55	6
ATOM	1767	CB	PRO A		7.9				1.00	20.33	
							32.893	60.702			6
ATOM	1768	CG	PRO A		8.6	26	33.068	62.046	1.00	31.02	6
ATOM	1769	С	PRO A	223	9.6	45	34.205	59.366	1.00	29.20	6
ATOM	1770	0	PRO A	223	10.69	94	34.680	59.796	1.00		8
MOTA	1771	N	LYS A	224	8.8		34.841	58.521	1.00		
											7
ATOM	1772	CA	LYS A		9.13		36.172	58.026	1.00		6
MOTA	1773	CB	LYS A		8.28	35	36.443	56.766	1.00	24:71	6
ATOM	1774	CG	LYS A	224	8.56		35.500	55.619	1.00 2		6
ATOM	1775	CD	LYS A		7.73			54.394	1.00		
							35.800				6
MOTA	1776	CE	LYS A		8.06		34.769	53.329	1.00		6
MOTA	1777	NZ	LYS A		7.19	8	34.849	52.122	1.00		7
ATOM	1778	С	LYS A	224	8.70		37.151	59.111	1.00 2		6
ATOM	1779	Ō	LYS A		7.99		36.780	60.055	1.00	22 62	8
			GLY A								
ATOM	1780	N			9.12		38.404	58.960	1.00 2		7
atom	1781	CA	GLY A		8.77		39.432	59.925	1.00 2	29.80	6
ATOM	1782	С	GLY A	225	9.39	6	39.188	61.286	1.00 3	12.96	6
-				٠.		-		-	_		

MOTA	1783	O GLY A 225	9.068	39.861	62.271	1.00 31.20	8
ATOM	1784	N LEU A 226	10.299	38.216	61.338	1.00 32.86	7
							6
MOTA	1785	CA LEU A 226	10.975	37.877	62.575	1.00 34.55	
ATOM	1786	CB LEU A 226	12.149	36.958	62.255	1.00 34.46	6
MOTA	1787	CG LEU A 226	12.982	36.413	63.407	1.00 34.48	6
ATOM	1788	CD1 LEU A 226	12.146	35.425	64.212	1.00 33.18	6
ATOM	1789	CD2 LEU A 226	14.207	35.724	62.847	1.00 31.39	6
ATOM	1790	C LEU A 226	11.481	39.160	63.255	1.00 36.29	6
	1791	O LEU A 226	12.156	39.970	62.613	1.00 33.87	8
ATOM							
ATOM	1792	N ASN A 227	11.131	39.358	64.531	1.00 37.31	7
MOTA	1793	CA ASN A 227	- 11.592	40.536	65.279	1.00 37.26	6
ATOM	1794	CB ASN A 227	10.444	41.212	66.053	1.00 35.57	6
MOTA	1795	CG ASN A 227	9.920	40.368	67.208	1.00 36.07	6
ATOM	1796	OD1 ASN A 227	10.678	39.940	68.089	1.00 35.08	8
ATOM	1797	ND2 ASN A 227	8.611	40.143	67.218	1.00 32.33	7
ATOM	1798	C ASN A 227	12.688	40.096	66.250	1.00 37.95	6
	1799	O ASN A 227	12.869	38.890	66.473	1.00 37.08	-8
ATOM							7
ATOM	1800	N ASP A 228	13.403	41.063	66.832	1.00 36.07	
ATOM	1801	CA ASP A 228	14.505	40.754	67.751	1.00 37.63	6
MOTA	1802	CB ASP A 228	14.996	42.007	68.486	1.00 36.48	6
ATOM	1803	CG ASP A 228	15.480	43.088	67.545	1.00 37.52	6
MOTA	1804	OD1 ASP A 228	15.936	42.752	66.427	1.00 35.28	8
ATOM	1805	OD2 ASP A 228	15.426	44.274	67.937	1.00 39.01	8
ATOM	1806	C ASP A 228	14.204	39.678	68.783	1.00 37.56	6
ATOM	1807	O ASP A 228	14.921	38.678	68.869	1.00 39.53	8
	1808	N ASN A 229	13.155	39.889	69.572	1.00 38.37	7
MOTA		CA ASN A 229	12.766		70.605	1.00 37.49	6
ATOM	1809			38.935	71.200	1.00 37.49	
MOTA	1810	CB ASN A 229	11.422	39.352			6
MOTA	1811	CG ASN A 229	11.490	40.709	71.877	1.00 40.47	6
ATOM	1812	OD1 ASN A 229	12.041	40.840	72.973	1.00 41.76	8
MOTA	1813	ND2 ASN A 229	10.960	41.735	71.212	1.00 36.50	7
MOTA	1814	C ASN A 229	12.680	37.530	70.017	1.00 37.64	6
ATOM	1815	O ASN A 229	13.446	36.634	70.395	1.00 35.76	. 8
MOTA	1816	N GLU A 230	11.758	37.351	69.076	1.00 36.01	7
ATOM	1817	CA GLU A 230	11.574	36.062	68.425	1.00 34.74	6
MOTA	1818	CB GLU A 230	10.753	36.242	67.153	1.00 35.55	6
	1819	CG GLU A 230	9.382	36.820	67.407	1.00 36.95	6
ATOM			8.580	36.960	66.144	1.00 35.30	6
ATOM	1820						
MOTA	1821	OE1 GLU A 230	9.042	37.670	65.229	1.00 36.98	8
ATOM	1822	OE2 GLU A 230	7.490	36.361	66.065	1.00 36.71	8
ATOM	1823	C GLU A 230	12.916	35.421	68.082	1.00 33.92	6
MOTA	1824	O GLU A 230	13.143	34.238	68.346	1.00 32.74	8
ATOM	1825	N PHE A 231	13.804	36.207	67.487	1.00 32.03	7
ATOM	1826	CA PHE A 231	15.116	35.712	67.123.	1.00 30.55	6
ATOM	1827	CB PHE A 231	15.932	36.821	66.460	1.00 33.86	6
ATOM	1828	CG 'HE A 231	17.295	36.381	66.012	1.00 36.97	` 6
ATOM	1829	CD1 -HE A 231	17.438	35.334	65.102	1.00 40.41	6
ATOM	1830	CD2 .HE A 231	18.436	37.021	66.480	1.00 36.58	6
•				34.932	64.661	1.00 43.00	6
MOTA	1831		18.709				
MOTA	1832	CE2 PHE A 231	19.711	36.632	66.049	1.00 39.07	6
MOTA	1833	CZ PHE A 231	19.849	35.586	65.137	1.00 40.52	6
ATOM	1834	C PHE A 231	15.835	35.232	68.376	1.00 30.63	6
MOTA	1835	O PHE A 231	16.177	34.042	68.497	1.00 29.66	8
ATOM	1836	N LEU A 232	16.049	36.162	69.310	1.00 24.94	7
ATOM	1837	CA LEU A 232	16.742	35.857	70.556	1.00 22.82	6
ATOM	1838	CB LEU A 232	16.724	37.084	71.468	1.00 24.96	. 6
	1839	CG LEU A 232	17.507	38.282	70.890	1.00 29.34	6
MOTA	•	CD1 LEU A 232	17.316	39.549	71.746	1.00 24.38	6
MOTA	1840				70.787	1.00 27.39	6
MOTA	1841	CD2 LEU A 232	18.991	37.903			
MOTA	1842	C LEU A 232	16.150	34.638	71.261	1.00 22.44	6
MOTA	1843	O LEU A 232	16.882	33.793	71.767	1.00 20.37	8
MOTA	1844	N PHE A 233	14.825	34.552	71.289	1.00 24.81	7
ATOM	1845	CA PHE A 233	14.131	33.422	71.905	1.00 25.81	6
MOTA	1846	CB PHE A 233	12.623	33.535	71.641	1.00 24.37	6
ATOM	1847	CG PHE A 233	11.811	32.373	72.157	1.00 24.18	6
ATOM	1848	CD1 PHE A 233	11.491	32.264	73.503	1.00 25.59	6

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ATOM 1849 CD2 PHE A 23	3 1	1.339	31.397	71.284	1 00 25	25
ATOM 1850 CE1 PHE A 23	3 1		31.198	73.974	1.00 25 1.00 25	
ATOM 1851 CE2 PHE A 23	3.1		30.327	71.747	1.00 25	
ATOM 1852 CZ PHE A 23		_	30.232	73.093	1.00 22	
ATOM 1853 C PHE A 23		4.661 3	32.133	71.275	1.00 26	.64
ATOM- 1854 O PHE A 23		5.094 3	31.209	71.971	1.00 28	
ATOM 1855 N ALA A 23		4.624 3	32.087	69.949	1.00 27	.27
ATOM 1856 CA ALA A 234 ATOM 1857 CB ALA A 234			30.921	69.209	1.00 30	.24
ATOM 1857 CB ALA A 234 ATOM 1858 C ALA A 234		4.797 3	31.107	67.720	1.00 33	
ATOM 1859 O ALA A 234			0.645	69.433	1.00 30	.76 6
ATOM 1860 N LEU A 235	_		9.491	69.488	1.00 30	.04 8
ATOM 1861 CA LEU A 235		_	31.695 31.486	69.563	1.00 31.	
ATOM 1862 CB LEU A 235			2.819	69.790-	1.00 32.	.83 6
ATOM 1863 CG LEU A 235			2.745	69.703 69.316	1.00 34.	
ATOM 1864 CD1 LEU A 235	21			69.205	1.00 36. 1.00 36.	
ATOM 1865 CD2 LEU A 235	21	:		70.330	1.00 35.	
ATOM 1866 C LEU A 235	18			71.176	1.00 30.	
ATOM 1867 O LEU A 235	19		_	71.312	1.00 30.	12 8
ATOM 1868 N GLU A 236	18			72.192	1.00 29.	
ATOM 1869 CA GLU A 236		1.418 3		73.561	1.00 33.	
ATOM 1870 CB GLU A 236				74.452	1.00 35.	06 6
ATOM 1871 CG GLU A 236 ATOM 1872 CD GLU A 236		.843 33	3.176	74.635	1.00 42.	35 6
ATOM 1872 CD GLU A 236 ATOM 1873 OE1 GLU A 236				74.843	1.00 47.	
ATOM 1874 OE2 GLU A 236				75.556	1.00 48.	
ATOM 1875 C GLU A 236				74.297	1.00 49.	
ATOM 1876 O GLU A 236			_	73.639	1.00 34.	
ATOM 1877 N LYS A 237	16			74.116 73.176	1.00 30.	
ATOM 1878 CA LYS A 237	16			73.175	1.00 35.	
ATOM 1879 CB LYS A 237			-	72.452	1.00 37.	
ATOM 1880 CG LYS A 237			_	72.848	1.00 39.	
ATOM 1881 CD LYS A 237				73.821	1.00 40.	66 6
ATOM 1882 CE LYS A 237				75.153	1.00 40.5	
ATOM 1883 NZ LYS A 237 ATOM 1884 C LYS A 237				76.083	1.00 32.0	05 7
ATOM 1885 O LYS A 237				72.485	1.00 35.0	
ATOM 1886 N SER A 238				73.061	1.00 30.6	
ATOM 1887 CA SER A 238					1.00 33.5	
ATOM 1888 CB SER A 238					1.00 32.1 1.00 32.0	
ATOM 1889 OG SER A 238					1.00 32.0	
ATOM 1890 C SER A 238					1.00 31.9	
ATOM 1891 O SER A 238	20.	064 24			1.00 26.8	
ATOM 1892 N LEU A 239			.920 7		1.00 31.1	
ATOM 1893 CA LEU A 239				2.147	1.00 30.8	
ATOM 1894 CB LEU A 239 ATOM 1895 CG LEU A 239	22.		.914 7		1.00 28.0	
ATOM 1896 CD1 LEU A 239	22. 23.				1.00 24.3	
ATOM 1897 CD2 LEU A 239	23.				1.00 24.6	
ATOM 1898 C LEU A 239	21.				1.00 23.1 1.00 31.9	
ATOM 1899 O LEU A 239	22.				1.00 31.9	
ATOM 1900 N GLU A 240	20.:				1.00 32.3	. •
ATOM 1901 CA GLU A 240	20.0				1.00 38.6	
ATOM 1902 CB GLU A 240	18.7				L.00 43.2	
ATOM 1903 CG GLU A 240	18.5		468 77	7.045 1	1.00 53.5	2 6
ATOM 1904 CD GLU A 240	19.6		383 78	3.022 1	00 56.39	
ATOM 1905 OE1 GLU A 240	19.9		399 78	3.701 1	00 57.8	18
ATOM 1906 OE2 GLU A 240 ATOM 1907 C GLU A 240	20.3				00 55.1	2 8
ATOM 1907 C GLU A 240 ATOM 1908 O GLU A 240	20.0	•			00 39.28	
ATOM 1909 N ILE A 241	20.5 19.4				.00 38.83	
ATOM 1910 CA ILE A 241	19.4				00 40.74	
ATOM 1911 CB ILE A 241	18.4				.00 38.08 .00 33.57	
ATCM - 1912 CG2 ILE A 241	18.5				.00 33.37	
ATOM 1913 CG1 ILE A 241	17.0			.056 1	.00 31.06	5 6
ATOM 1914 CD1 ILE A 241	16.1			.843 1	.00 27.53	6
			-		,	

ATOM	1919	5 C	ILE A	241	2	0.713	21.3	72 72.74	7 1.00	39.56	6
ATOM	1916	5 0	ILE A	241	2	0.984	20.18	89 72.93		40.82	
· ATOM	1917		VAL A			1.605		54 72.29	9 1.00	41.93	
MOTA	1918					2.979				45.09	
MOTA	1919					3.808				45.76	_
ATOM	1920 1921		31 VAL A 32 VAL A			5.242				43.09	
MOTA MOTA	1921		VAL A			3.182				46.41	
ATOM	1923		VAL A			3.698 1.191				45.69 46.30	_
ATOM	1924		LYS A			3.750				46.30	
ATOM	1925					1.427				46.96	6
ATOM	1926					.214				49.49	6
MOTA	1927	CG	LYS A	243		.061				54.90	6
MOTA					24	1.652	23.93			58.95	6
ATOM	1929					.782	25.39	9 78.57	7 1.00	64.13	6
MOTA	1930					.274	26.28			66.93	7
ATOM	1931 1932		LYS A			.965	20.76			47.06	6
MOTA MOTA	1932	И	LYS A GLU A			.735	20.11			46.39	8
MOTA	1934	CA				.172	20.38 19.13			47.51	. 7 . 6
MOTA	1935	CB				.650	19.06			54.49	6
ATOM	19¦36	CG				.843	20.19			62.61	6
ATOM	1937	CD			18	.360	20.08			65.15	6
MOTA	1938		1 GLU A			.572	20.98	0 76.88	B · 1.00	66.49	8
ATOM	1939	OE:				.986	19.10			64.82	8
ATOM	1940 1941	C	GLU A :			.745	17.93			50.17	6
ATOM ATOM	1941	O N	VAL A			.866	16.84 18.14			51.54	8
MOTA	1943	CA	VAL A			.587	17.06			47.70 45.43	7 6
ATOM	1944	CB	VAL A			.704	16.98			48.47	6
MOTA	1945	CG:	l VAL A :	245		.082	15.76			51.87	6
ATOM	1946		2 VAL A		21	.226	16.93	4 72.731	L 1.00	45.65	6
ATOM	1947	C	VAL A			.056	17.07			43.01	6
ATOM	1948	0	VAL A			.620	16.00			39.28	8
ATOM ATOM	1949 1950	N CA	PHE A 2			.682 .063	18.24 18.32			40.53 38.56	7
ATOM	1951	CB	PHE A 2			.023	18.70			36.85	6 6
ATOM	1952	CG	PHE A 2			.315	18.48			36.46	6
MOTA	1953	CD1	PHE A 2	246		.749	17.20			32.95	6
ATOM	1954		PHE A 2		29	.064	19.58			35.51	6
ATOM	1955	CE1				.903	17.00	-		33.80	6
MOTA	1956 1957	CE2	PHE A 2 PHE A 2			.222	19.39	7 69.206		34.46	6
ATOM ATOM	1958	C	PHE A 2			.640 .970	18.103 19.313			35.54 40.83	6 6
ATOM	1959	ō	PHE A 2			613	20.478			40.32	8
ATOM	1960	N	GLU A 2			141	18.839			42.54	7
ATOM	1961	CA	GLU A 2	47	30.	128	19.695			43.93	6
ATOM	1962	CB	GLU A 2			655	19.075			45.67	6
ATOM	1963	CG	GLU A 2			763	19.243			51.63	6
ATOM	1964	CD OF1	GLU A 2			478	18.424			57.42	6
MOTA MOTA	1965 1966	OE1	GLU A 2 GLU A 2			645 296	18.644 17.557			62.12 59.43	8
ATOM	1967	c	GLU A 2			268	19.839			43.62	8 6
ATOM ·	1968	ō	GLU A 2			077	18.931			44.25	8
ATOM	1969	N	PRO A 2			342	20.988			43.65	7
ATOM	1970	CD	PRO, A 2	48	30.	439	22.143			42.73	6
ATOM	1971	CA	PRO A 2		32.		21.260	71.779	1.00	43.28	6
MOTA	1972	CB	PRO A 2		31.		22.480			43.03	6
ATOM	1973	CG	PRO A 2		30.		22.474	71.415		43.02	6
ATOM	1974	C	PRO A 24		33.		21.552			43.37	6
atom atom	1975 1976	O N	GLU A 24		33. 34.		22.286			45.44 42.38	8 7
ATOM	1977	CA	GLU A 24		36.		21.263			42.36	6
ATOM	1978	CB	GLU A 24		37.		20.275			42.06	6
ATOM	1979	CG	GLU A 24		36.		18.816			44.28	6
ATOM	1980	CD	GLU A 24		38.	015	17.908		1.00	44.79	6
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ATOM 1992 N TYR A 251 34.539 25.810 68.690 1.00 29.71 7 ATOM 1993 CA TYR A 251 32.86 26.451 68.860 1.00 27.07 6 ATOM 1995 CG TYR A 251 32.080 27.872 69.406 1.00 21.01 6 ATOM 1995 CG TYR A 251 32.080 27.872 69.406 1.00 31.46 6 ATOM 1996 CD1 TYR A 251 31.553 28.903 68.622 1.00 31.46 6 ATOM 1997 CEL TYR A 251 31.439 01.90 69.106 1.00 31.65 6 ATOM 1998 CD2 TYR A 251 32.494 28.181 70.696 1.00 31.65 6 ATOM 1999 CEZ TYR A 251 32.494 28.181 70.696 1.00 31.45 6 ATOM 1999 CTZ TYR A 251 31.439 01.90 69.106 1.00 31.55 6 ATOM 2001 CZ TYR A 251 31.439 10.196 69.106 1.00 33.89 6 ATOM 2001 OH TYR A 251 31.431 31.773 70.867 1.00 33.89 6 ATOM 2002 C TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2003 O TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2004 N LEU A 252 33.631 27.73 66.3451 1.00 24.78 8 ATOM 2005 CA LEU A 252 33.631 27.810 63.451 1.00 24.78 8 ATOM 2006 CB LEU A 252 33.631 27.810 63.451 1.00 21.32 6 ATOM 2007 CG LEU A 252 33.631 27.810 63.451 1.00 21.32 6 ATOM 2008 CD1 LEU A 252 33.631 27.810 63.451 1.00 22.50 6 ATOM 2001 C LEU A 252 33.632 64.815 1.00 21.32 6 ATOM 2001 C LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2010 C LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2010 C LEU A 253 35.497 26.399 61.986 1.00 22.07 6 ATOM 2011 O LEU A 253 35.499 26.999 61.986 1.00 22.07 6 ATOM 2010 C LEU A 253 35.495 26.999 61.986 1.00 22.07 6 ATOM 2011 C LEU A 253 35.495 26.999 61.986 1.00 22.07 6 ATOM 2012 N LEU A 253 30.352 31.794 65.595 1.00 22.55 6 ATOM 2013 CA LEU A 253 33.734 67.29 1.00 22.56 6 ATOM 2016 CD1 LEU A 253 30.352 31.944 65.695 1.00 22.90 6 ATOM 2017 CD2 LEU A 253 33.734 30.277 64.554 1.00 21.01 7 ATOM 2018 C LEU A 253 30.352 31.944 65.695 1.00 22.90 6 ATOM 2010 C LEU A 253 30.352 31.944 65.695 1.00 22.90 6 ATOM 2010 C LEU A 253 30.353 31.734 30.277 64.554 1.00 22.90 6 ATOM 2013 CA LEU A 253 30.356 61.856 61.869 1.00 22.91 6 ATOM 2020 N GLEU A 253 30.356 61.856 61.869 1.00 22.93 6 ATOM 2021 CA GLEU A 253 30.356 61.586 82.997 1.00 18.17 6 ATOM 2022 CB GLEU A 255 29.966 31.755 62.864 1.00 22.89 6 ATOM 2030	MOTA	1991	. 0	VAL A 250	34.162	23.610	68.421	1.00 33.96	8
ATOM 1994 CB TYR A 251 33.368 26.183 67.916 1.00 27.07 6 ATOM 1995 CG TYR A 251 32.080 27.872 68.406 1.00 31.46 6 ATOM 1996 CD1 TYR A 251 31.533 28.903 68.622 1.00 31.46 6 ATOM 1996 CD1 TYR A 251 31.439 30.196 69.106 1.00 31.46 6 ATOM 1998 CD2 TYR A 251 31.439 30.196 69.106 1.00 31.46 6 ATOM 1998 CD2 TYR A 251 32.494 28.181 70.896 1.00 30.20 6 ATOM 2000 CZ TYR A 251 31.439 30.196 69.106 1.00 30.20 6 ATOM 2000 CZ TYR A 251 31.437 30.492 77.71.193 1.00 34.75 6 ATOM 2001 OH TYR A 251 31.474 31.773 70.896 71.00 33.89 6 ATOM 2001 OH TYR A 251 31.474 31.773 70.896 71.00 33.52 8 ATOM 2001 OH TYR A 251 31.474 31.773 70.896 71.00 33.52 8 ATOM 2003 O TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2000 CC TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2003 O TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2003 O TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2003 O TYR A 251 33.570 27.384 66.992 1.00 27.48 6 ATOM 2005 CA LEU A 252 33.163 27.254 65.773 1.00 24.80 7 ATOM 2005 CA LEU A 252 33.150 28.332 64.815 1.00 21.32 6 ATOM 2005 CA LEU A 252 33.631 27.810 63.451 1.00 21.32 6 ATOM 2007 CG LEU A 252 33.457 26.373 64.395 1.00 22.51 6 ATOM 2008 CD1 LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2001 O LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2010 C LEU A 252 31.752 28.266 64.856 1.00 21.99 8 ATOM 2011 O LEU A 253 31.750 28.266 64.856 1.00 21.99 7 ATOM 2012 N LEU A 253 31.750 28.266 64.856 1.00 21.01 7 ATOM 2013 CA LEU A 253 31.750 28.266 64.856 1.00 21.01 7 ATOM 2012 CA LEU A 253 30.530 27.254 65.842 1.00 21.01 7 ATOM 2013 CA LEU A 253 31.750 28.266 66 64.856 1.00 21.01 7 ATOM 2015 CG LEU A 253 30.539 31.91 67.145 1.00 18.89 6 64.00 18.8				TYR A 251					
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ATOM 1996 CD1 TYR A 251 31.553 28.903 68.622 1.00 31.14 6 ATOM 1998 CD2 TYR A 251 31.439 30.196 69.106 1.00 31.66 6 ATOM 1998 CD2 TYR A 251 32.494 28.181 70.696 1.00 30.20 6 ATOM 1999 CE2 TYR A 251 32.384 29.477 71.193 1.00 33.89 6 ATOM 2000 CZ TYR A 251 31.854 30.482 70.391 1.00 34.75 6 ATOM 2001 OH TYR A 251 31.743 31.773 70.867 1.00 33.52 6 ATOM 2002 C TYR A 251 33.570 27.384 66.922 1.00 27.48 6 ATOM 2003 O TYR A 251 33.570 27.384 66.922 1.00 27.48 6 ATOM 2003 O TYR A 251 33.570 27.384 66.922 1.00 27.48 6 ATOM 2005 CA LEU A 252 33.630 27.254 65.773 1.00 24.78 8 ATOM 2005 CA LEU A 252 33.150 28.332 64.815 1.00 24.78 8 ATOM 2006 CB LEU A 252 33.150 28.332 64.815 1.00 23.40 6 ATOM 2007 CG LEU A 252 33.150 28.332 64.815 1.00 21.84 6 ATOM 2008 CD1 LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2009 CD2 LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2010 C LEU A 252 31.762 28.999 61.986 1.00 22.56 6 ATOM 2011 O LEU A 252 31.762 28.999 64.729 1.00 22.56 6 ATOM 2012 N LEU A 253 31.754 30.277 64.554 1.00 21.01 7 ATOM 2013 CA LEU A 253 31.754 30.277 64.554 1.00 21.01 7 ATOM 2013 CA LEU A 253 31.754 30.277 64.554 1.00 21.01 7 ATOM 2014 CB LEU A 253 30.530 31.901 63.198 1.00 22.56 6 ATOM 2015 CG LEU A 253 30.530 31.901 63.198 1.00 22.56 6 ATOM 2016 CD1 LEU A 253 30.530 31.904 65.654 1.00 21.01 7 ATOM 2017 CD2 LEU A 253 30.530 31.901 63.198 1.00 22.56 6 ATOM 2018 C LEU A 253 30.530 31.901 63.198 1.00 22.56 6 ATOM 2015 CG LEU A 253 30.530 31.901 63.998 1.00 22.55 6 ATOM 2016 CD1 LEU A 253 30.530 31.901 63.198 1.00 22.55 6 ATOM 2017 CD2 LEU A 253 30.530 31.901 63.198 1.00 22.00 6 ATOM 2018 C LEU A 253 31.466 32.691 62.987 1.00 18.17 8 ATOM 2020 N GLN A 254 29.849 32.906 61.00 2.200 6 ATOM 2018 C LEU A 253 31.466 32.691 62.987 1.00 18.17 8 ATOM 2020 N GLN A 254 29.849 32.906 61.00 2.302 7 ATOM 2021 CA GLN A 254 29.848 32.906 61.00 2.302 7 ATOM 2022 CB GLN A 254 29.848 32.906 61.007 1.00 18.17 6 ATOM 2023 CG LEU A 255 28.905 34.854 61.007 1.00 19.40 7 ATOM 2024 CB LEU A 255 28.905 36.506 60.149 1.00 22.90 6 ATOM 2035 CG	АТОМ	1995	CG	TYR A 251	32.080	27.872	69.406	1.00 31.46	6
ATOM						_			
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ATOM 2008 CD1 LEU A 252 35.457 26.373 64.395 1.00 22.51 6 ATOM 2010 C LEU A 252 35.499 26.999 61.986 1.00 22.56 6 ATOM 2011 O LEU A 252 31.762 28.959 64.729 1.00 22.55 6 ATOM 2012 N LEU A 253 31.734 30.277 64.554 1.00 21.99 8 ATOM 2013 CA LEU A 253 31.734 30.277 64.554 1.00 21.01 7 ATOM 2014 CB LEU A 253 30.498 31.047 64.461 1.00 18.89 6 ATOM 2015 CG LEU A 253 29.198 32.942 65.842 1.00 21.61 6 ATOM 2016 CD1 LEU A 253 29.198 32.942 65.842 1.00 21.61 6 ATOM 2017 CD2 LEU A 253 29.395 33.716 67.145 1.00 22.90 6 ATOM 2018 C LEU A 253 30.539 31.904 63.198 1.00 20.05 6 ATOM 2019 O LEU A 253 29.395 33.716 67.145 1.00 22.90 6 ATOM 2019 C LEU A 253 30.539 31.901 63.198 1.00 20.05 6 ATOM 2020 N GLN A 254 29.544 31.720 62.987 1.00 18.17 8 ATOM 2021 CA GLN A 254 29.544 31.720 63.40 1.00 19.40 7 ATOM 2022 CB GLN A 254 29.548 32.490 61.115 1.00 18.17 8 ATOM 2022 CB GLN A 254 29.917 31.592 59.969 1.00 9.67 6 ATOM 2023 CG GLN A 254 27.584 31.713 59.601 1.00 19.97 6 ATOM 2025 CE GLN A 254 27.584 31.713 59.601 1.00 18.97 6 ATOM 2027 C GLN A 254 27.584 31.713 59.601 1.00 19.97 6 ATOM 2028 O GLN A 254 27.584 31.713 59.601 1.00 18.97 6 ATOM 2029 N LEU A 255 28.905 33.634 61.444 1.00 19.75 6 ATOM 2029 N LEU A 255 28.905 33.634 61.444 1.00 19.75 6 ATOM 2027 C GLN A 254 27.584 31.713 59.601 1.00 22.54 8 ATOM 2028 CB LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2027 C GLN A 254 26.564 33.769 58.869 1.00 22.77 8 ATOM 2028 CB LEU A 255 28.936 36.993 62.242 1.00 23.77 6 ATOM 2030 CA LEU A 255 28.936 36.993 62.242 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.936 36.993 62.242 1.00 23.77 6 ATOM 2033 CB LEU A 255 28.936 36.993 62.242 1.00 23.77 6 ATOM 2034 CB LEU A 255 27.974 38.066 60.149 1.00 29.34 6 ATOM 2035 C LEU A 255 27.974 38.066 60.149 1.00 29.34 6 ATOM 2036 CA LEU A 255 27.974 38.066 60.149 1.00 29.28 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 23.77 1 6 ATOM 2038 CA GLY A 256 26.969 36.158 59.254 1.00 27.11 6 ATOM 2034 CB THR A 257 25.599 39.150 58.072 1.00 27.85 6 ATOM 2040 CB THR A 257 25.599 39.150 58.072 1.00 27.85 6 ATOM 2	MOTA	2007	CG	LEU A 252	35.126	27.456	63.385	1.00 21.84	6
ATOM 2010 CD2 LEU A 252 35.499 26.999 61.986 1.00 22.07 6 ATOM 2011 C LEU A 252 31.762 28.959 64.729 1.00 22.56 6 ATOM 2011 C LEU A 252 30.750 28.266 64.854 1.00 21.99 8 ATOM 2012 N LEU A 253 31.734 30.277 64.554 1.00 21.01 7 ATOM 2013 CA LEU A 253 30.498 31.047 64.461 1.00 18.89 6 ATOM 2014 CB LEU A 253 30.498 31.047 64.461 1.00 18.89 6 ATOM 2015 CG LEU A 253 29.198 32.942 65.842 1.00 21.61 6 ATOM 2016 CD1 LEU A 253 27.849 32.220 65.860 1.00 22.23 6 ATOM 2017 CD2 LEU A 253 29.395 33.716 67.145 1.00 22.23 6 ATOM 2018 C LEU A 253 30.539 31.901 63.198 1.00 20.05 6 ATOM 2019 O LEU A 253 31.466 32.691 62.987 1.00 18.17 8 ATOM 2020 N GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2021 CA GLN A 254 29.488 32.490 61.115 1.00 18.17 8 ATOM 2022 CB GLN A 254 29.917 31.592 59.969 1.00 19.40 7 ATOM 2023 CG GLN A 254 29.017 31.592 59.969 1.00 19.40 7 ATOM 2024 CD GLN A 254 27.584 31.713 59.601 1.00 19.40 7 ATOM 2025 OE1 GLN A 254 27.584 31.713 59.601 1.00 19.40 7 ATOM 2027 C GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2027 C GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2027 C GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2027 C GLN A 254 27.917 32.677 57.450 1.00 22.89 7 ATOM 2028 O GLN A 254 27.917 32.677 57.450 1.00 22.89 7 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2031 CB LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2033 CD1 LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2031 CB LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2033 CD1 LEU A 255 27.902 36.506 64.456 1.00 29.34 6 ATOM 2036 CA LEU A 255 27.902 36.506 64.456 1.00 29.34 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 23.02 7 ATOM 2038 CA GLY A 256 26.969 36.158 59.254 1.00 23.02 7 ATOM 2036 CA GLY A 256 26.969 36.158 59.254 1.00 23.02 7 ATOM 2036 CA GLY A 256 26.969 36.158 59.254 1.00 23.02 7 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.67 8 ATOM 2038 CA GLY A 256 26.969 36.158 59.254 1.00 27.88 6 ATOM 2034 CB THR A 257 25.599 39.150 58.072 1.00 27.85 6 ATO		_							
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ATOM 2012 N LEU A 253 31.734 30.277 64.554 1.00 21.01 7 ATOM 2013 CA LEU A 253 30.498 31.047 64.461 1.00 18.89 6 ATOM 2015 CG LEU A 253 20.492 65.860 1.00 21.61 6 ATOM 2016 CD1 LEU A 253 27.849 32.220 65.860 1.00 22.23 6 ATOM 2017 CD2 LEU A 253 27.849 32.220 65.860 1.00 22.23 6 ATOM 2018 C LEU A 253 30.539 31.901 63.198 1.00 20.05 6 ATOM 2019 O LEU A 253 31.466 32.691 62.987 1.00 18.17 8 ATOM 2020 N GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2022 CB GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2022 CB GLN A 254 29.017 31.592 59.969 1.00 9.67 6 ATOM 2023 CG GLN A 254 27.584 31.713 59.601 1.00 18.17 6 ATOM 2026 NE2 GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2026 NE2 GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2027 C GLN A 254 28.520 33.634 61.444 1.00 19.75 6 ATOM 2028 O GLN A 254 28.520 33.634 61.444 1.00 19.75 6 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2031 CB LEU A 255 28.905 36.856 61.369 1.00 29.34 6 ATOM 2031 CB LEU A 255 28.905 36.856 61.369 1.00 29.34 6 ATOM 2032 CG GLU A 255 28.905 36.856 61.369 1.00 29.34 6 ATOM 2031 CB LEU A 255 28.905 36.856 61.369 1.00 29.34 6 ATOM 2032 CG LEU A 255 28.905 36.856 61.369 1.00 29.34 6 ATOM 2031 CB LEU A 255 28.905 36.856 61.369 1.00 28.42 6 ATOM 2033 CD LEU A 255 28.905 36.856 61.369 1.00 28.42 6 ATOM 2034 CD LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 CA GLY A 256 26.969 36.158 59.254 1.00 27.85 7 ATOM 2038 CA GLY A 256 26.969 36.158 59.254 1.00 27.85 7 ATOM 2039 C GLY A 256 26.969 36.158 59.254 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2042 CA THR A 257 25.599 39.150 58.604 1.00 27.98 6 ATOM 2045 CG2 THR A 257 25.599 39.150 58.604 1.00 27.98 6 ATOM 2045 CG2 THR A 257 25.599 39.150 58.505 1.00 27.98 6 ATOM 2045 CG2 THR A 257 25.599 39.150 58.505 1.00 27.98 6 ATOM 2045 CG2 THR A 257 25.599 39.150 58.000 31.91 8 ATOM 2045 CG2 THR A 257 26.606 41.541 59.510 1.00 26.45 6	ATOM	2011	0	LEU A 252	30.750	28,266	64.856	1.00 21.99	8
ATOM 2013 CA LEU A 253 30.498 31.047 64.461 1.00 18.89 6 ATOM 2014 CB LEU A 253 30.352 31.944 65.695 1.00 20.05 6 ATOM 2015 CG LEU A 253 29.198 32.942 65.842 1.00 21.61 6 ATOM 2016 CD1 LEU A 253 29.395 33.716 67.145 1.00 22.23 6 ATOM 2017 CD2 LEU A 253 29.395 33.716 67.145 1.00 22.90 6 ATOM 2018 C LEU A 253 30.539 31.901 63.198 1.00 20.05 6 ATOM 2019 O LEU A 253 31.466 32.691 62.987 1.00 18.17 8 ATOM 2020 N GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2021 CA GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2022 CB GLN A 254 29.017 31.592 59.699 1.00 9.67 6 ATOM 2023 CG GLN A 254 27.584 31.713 59.601 1.00 18.43 6 ATOM 2024 CD GLN A 254 27.584 31.713 59.601 1.00 18.43 6 ATOM 2025 OE1 GLN A 254 27.584 31.713 59.601 1.00 18.97 6 ATOM 2026 NE2 GLN A 254 27.584 31.713 59.601 1.00 22.54 8 ATOM 2026 NE2 GLN A 254 27.584 31.713 59.601 1.00 22.54 8 ATOM 2026 NE2 GLN A 254 27.584 31.713 59.601 1.00 22.54 8 ATOM 2027 C GLN A 254 27.917 32.677 57.450 1.00 22.54 8 ATOM 2028 O GLN A 254 26.564 33.769 58.869 1.00 22.54 8 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.07 7 ATOM 2030 CA LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2033 CD LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2033 CD LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2033 CD LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2033 CD LEU A 255 28.905 34.854 61.067 1.00 24.84 6 ATOM 2031 CB LEU A 255 27.902 36.506 64.456 1.00 29.34 6 ATOM 2033 CD LEU A 255 27.902 36.506 60.149 1.00 24.84 6 ATOM 2036 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2030 CA LEU A 255 25.599 39.150 58.672 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.672 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.545 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.545 1.00 27.85 6 ATOM 2045 CG2 THR A 257 26.500 41.595 59.510 1.00 26.									
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ATOM 2016 CD1 LEU A 253 27.849 32.220 65.860 1.00 22.23 6 ATOM 2017 CD2 LEU A 253 29.395 33.716 67.145 1.00 22.90 6 ATOM 2018 C LEU A 253 30.539 31.901 63.198 1.00 20.05 6 ATOM 2019 0 LEU A 253 31.466 32.691 63.198 1.00 18.17 8 ATOM 2020 N GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2021 CA GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2022 CB GLN A 254 29.017 31.592 59.969 1.00 9.67 6 ATOM 2023 CG GLN A 254 27.584 31.713 59.601 1.00 18.43 6 ATOM 2024 CD GLN A 254 27.584 31.713 59.601 1.00 19.97 6 ATOM 2025 OEI GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2025 OEI GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2026 NE2 GLN A 254 26.564 33.769 58.869 1.00 22.54 8 ATOM 2027 C GLN A 254 26.564 33.769 58.869 1.00 22.89 7 ATOM 2028 O GLN A 254 28.520 33.634 61.444 1.00 19.75 6 ATOM 2029 N LEU A 255 28.963 36.993 62.242 1.00 23.02 7 ATOM 2030 CA LEU A 255 28.963 36.993 62.242 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.993 62.242 1.00 26.84 6 ATOM 2032 CG LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 29.226 36.556 64.436 1.00 29.34 6 ATOM 2035 C LEU A 255 27.902 36.506 64.436 1.00 29.34 6 ATOM 2036 C LEU A 255 27.902 36.506 64.456 1.00 29.34 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 26.84 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 26.11 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 27.11 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 27.11 6 ATOM 2031 CB LEU A 255 27.505 36.842 60.197 1.00 24.84 6 ATOM 2039 C GLY A 256 26.969 36.158 59.254 1.00 27.11 6 ATOM 2034 CB THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2040 C GLY A 256 26.408 36.858 58.117 1.00 27.85 7 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CB THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2045 CB THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2040 CB THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2045 CB THR A 257 26.606 41.541 59.510 1.00 26.45 6	MOTA	2015	CG	LEU A 253	29.198	32.942	65.842	1.00 21.61.	6
ATOM 2017 CD2 LEU A 253 29.395 33.716 67.145 1.00 22.90 6 ATOM 2018 C LEU A 253 30.539 31.901 63.198 1.00 20.05 6 ATOM 2019 O LEU A 253 31.466 32.691 62.987 1.00 18.17 8 ATOM 2020 N GLN A 254 29.544 31.720 62.340 1.00 19.40 7 ATOM 2021 CA GLN A 254 29.488 32.490 61.115 1.00 18.17 6 ATOM 2022 CB GLN A 254 29.017 31.592 59.969 1.00 9.67 6 ATOM 2023 CG GLN A 254 27.584 31.713 59.601 1.00 18.43 6 ATOM 2024 CD GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2025 OE1 GLN A 254 27.368 32.766 58.549 1.00 19.97 6 ATOM 2026 NE2 GLN A 254 27.917 32.677 57.450 1.00 22.54 8 ATOM 2026 NE2 GLN A 254 26.564 33.769 58.869 1.00 22.89 7 ATOM 2027 C GLN A 254 27.470 33.415 62.060 1.00 18.77 8 ATOM 2028 O GLN A 255 28.503 36.346 61.444 1.00 19.75 6 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2031 CB LEU A 255 28.905 34.854 61.067 1.00 23.07 7 ATOM 2031 CB LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.993 62.242 1.00 26.84 6 ATOM 2033 CD1 LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD LEU A 255 27.902 36.556 64.331 1.00 30.65 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 29.34 6 ATOM 2035 C LEU A 255 27.902 36.506 60.149 1.00 24.84 6 ATOM 2036 O LEU A 255 27.902 36.506 60.149 1.00 24.84 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2039 C GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2039 C GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2039 C GLY A 256 26.969 36.158 59.254 1.00 25.67 8 ATOM 2040 O GLY A 256 26.969 39.150 58.072 1.00 27.85 7 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2043 CB THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2043 CB THR A 257 25.599 59.150 58.072 1.00 27.98 6 ATOM 2045 CG2 THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2045 CG2 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.002 41.895 57.232 1.00 31.91 8									
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ATOM 2025 OE1 GLN A 254 27.917 32.677 57.450 1.00 22.54 8 ATOM 2026 NE2 GLN A 254 26.564 33.769 58.869 1.00 22.89 7 ATOM 2027 C GLN A 254 28.520 33.634 61.444 1.00 19.75 6 ATOM 2028 O GLN A 254 27.470 33.415 62.060 1.00 18.77 8 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2030 CA LEU A 255 28.905 34.854 61.067 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.952 61.369 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.993 62.242 1.00 26.84 6 ATOM 2032 CG LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 30.196 37.520 64.331 1.00 30.65 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 O LEU A 255 27.774 38.066 60.149 1.00 24.94 8 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 24.94 8 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 27.11 6 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 27.11 6 ATOM 2040 O GLY A 256 26.408 36.858 58.117 1.00 27.85 7 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2043 CB THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2045 CG2 THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2045 CG2 THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6		2024	CD	GLN A 254	27.368		58.549	1.00 19.97	6
ATOM 2026 NE2 GLN A 254 26.564 33.769 58.869 1.00 22.89 7 ATOM 2027 C GLN A 254 28.520 33.634 61.444 1.00 19.75 6 ATOM 2028 O GLN A 254 27.470 33.415 62.060 1.00 18.77 8 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2030 CA LEU A 255 28.132 36.052 61.369 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.953 62.242 1.00 26.84 6 ATOM 2032 CG LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 O LEU A 255 27.774 38.066 60.149 1.00 24.94 8 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2040 O GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2043 CB THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2044 OG1 THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									
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ATOM 2028 O GLN A 254 27.470 33.415 62.060 1.00 18.77 8 ATOM 2029 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2030 CA LEU A 255 28.132 36.052 61.369 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.993 62.242 1.00 26.84 6 ATOM 2032 CG LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 30.196 37.520 64.331 1.00 30.65 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 O LEU A 255 27.774 38.066 60.149 1.00 24.94 8 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2040 O GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6	ATOM	2027	C	GLN A 254		33.634	61.444	1.00 19.75	6
ATOM 2039 N LEU A 255 28.905 34.854 61.067 1.00 23.02 7 ATOM 2030 CA LEU A 255 28.132 36.052 61.369 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.993 62.242 1.00 26.84 6 ATOM 2032 CG LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 30.196 37.520 64.331 1.00 30.65 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 O LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 25.599 39.150 58.072 1.00 27.98 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.686 41.541 59.510 1.00 26.45 6		2028	0	GLN A 254	27.470	33.415	62.060	1.00 18.77	8
ATOM 2030 CA LEU A 255 28.132 36.052 61.369 1.00 23.77 6 ATOM 2031 CB LEU A 255 28.963 36.993 62.242 1.00 26.84 6 ATOM 2032 CG LEU A 255 29.226 36.556 63.684 1.00 29.34 6 ATOM 2033 CD1 LEU A 255 30.196 37.520 64.331 1.00 30.65 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 O LEU A 255 27.774 38.066 60.149 1.00 24.94 8 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2039 C GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2043 CB THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2044 OG1 THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									
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ATOM 2033 CD1 LEU A 255 30.196 37.520 64.331 1.00 30.65 6 ATOM 2034 CD2 LEU A 255 27.902 36.506 64.456 1.00 28.42 6 ATOM 2035 C LEU A 255 27.605 36.842 60.197 1.00 24.84 6 ATOM 2036 O LEU A 255 27.774 38.066 60.149 1.00 24.94 8 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6	ΔΤΟΜ	0032	CG	LEU A 255	29.226	36.556	63.684	1.00 29.34	6
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ATOM 2036 O LEU A 255 27.774 38.066 60.149 1.00 24.94 8 ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6	MOTA	2035	С	LEU A 255	27.605	36.842	60.197	1.00 24.84	6
ATOM 2037 N GLY A 256 26.969 36.158 59.254 1.00 25.07 7 ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6		2036	0	LEU A 255	27.774	38.066	60.149	1.00 24.94	8
ATOM 2038 CA GLY A 256 26.408 36.858 58.117 1.00 26.11 6 ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									7
ATOM 2039 C GLY A 256 25.506 37.956 58.644 1.00 27.11 6 ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									
ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6	ATOM								Ь
ATOM 2040 O GLY A 256 24.742 37.734 59.584 1.00 25.67 8 ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6	ATOM	2039	C	GLY A 256	25.506		58.644	1.00 27.11	6
ATOM 2041 N THR A 257 25.599 39.150 58.072 1.00 27.85 7 ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									
ATOM 2042 CA THR A 257 24.757 40.244 58.536 1.00 29.28 6 ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									ī
ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6									<i>-</i>
ATOM 2043 CB THR A 257 25.517 41.597 58.545 1.00 27.98 6 ATOM 2044 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6	MOTA					40.244			9
ATOM 2344 OG1 THR A 257 26.002 41.895 57.232 1.00 31.91 8 ATOM 2345 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6		2043	CB	THR A 257	25.517	41.597	58.545	1.00 27.98	6
ATOM 2045 CG2 THR A 257 26.686 41.541 59.510 1.00 26.45 6				THR A 257			57.232		8
7.01									É
ATOM 2046 C THR A 257 23.477 40.392 57.722 1.00 28.39 6									
	ATOM	2046	С	THR A 257	23.477	40.392	57.722	1.00 48.39	,

ATOM	2047	7 0	THR A	257		22.74	7	41.370	E 7	.879	1 0		40	_
ATOM			ASP A									29.		8
						23.19		39.414		.867		29.		7
MOTA						21.97		39.471		.065		30.	. 49	6
ATOM						22.00		38.432	54	.933		28,	. 22	6
ATOM	2051					22.33	7 3	37.033	55	.416	1.00	29.	.39	6
ATOM	2052	OI	1 ASP A	258		21.89	3 3	36.653	56	.520		30.		8
ATOM	2053		2 ASP A			23.01	-	6.292		. 667		29.		
ATOM	2054		ASP A			20.64		9.355						8
ATOM	2055		ASP A							.826		30.		6
ATOM	2056					19.60		9.622		.248		32.		8
			PRO A			20.65		8.912		.101		30.	-	7
ATOM	2057					21.71	_	8.338	58.	. 952	1.00	33.	56	6
ATOM	2058					19.36	63	8.821	58.	806	1.00	31.	73	6
ATOM	2059		PRO A	259		19.70	5 3	7.912	59.	987	1.00	31.	87	6
ATOM	2060	CG	PRO A	259		21.06	7 3	8.373	60.	333		31.		6
MOTA	2061	С	PRO A	259		18.817		0.184		260		30.		6
MOTA	2062	0	PRO A			17.736		0.270		84-5		29.		
ATOM	2063	N	LEU A			19.565		1.245		980				8
ATOM	2064	CA	LEU A									29.		7
ATOM	2065	CB	LEU A			19.161		2.592		375		29.		6
						20.367		3.542		275		27.		6
MOTA	2066	CG	LEU A			21.543		3.286		234		22.		6
ATOM	2067		L LEU A			22.794		3.919	59.	684	1.00	15.	79	6
ATOM	2068		LEU A		2	21.211	. 4	3.793	61.	633	1.00	16.	25	6
MOTA	2069	С	LEU A	260	1	L7.992	4	3.165	58.	576	1.00	28.	09	6
ATOM	2070	0	LEU A	260	1	17.787	4	2.834		410		29.		8
ATOM	2071	N	LEU A	261	1	7.237		4.044		223		29.		7
ATOM	2072	CA	LEU A	261		6.097		4.693		596		29.		6
ATOM	2073	CB	LEU A			5.540		5.788		513		29.		
ATOM	2074	CG	LEU A			4.406		6.664						6
ATOM	2075		LEU A							950		28.		6
ATOM	2076		LEU A			3.144		5.819		803		24.		6
						4.139	-	7.859	59.			25.		6
MOTA	2077	C	LEU. A			6.461		5.322	57.			29.		6
ATOM	2078	0	LEU A			5.717		5.198	56.	295	1.00	31.	67	8
MOTA	2079	N	GLU A		1	7.603	4!	5.998	57.	201	1.00	31.	54	7
MOTA	2080	,CA	GLU A		1	8.015	4 (5.664	55.	973	1.00	31.5	93	6
ATOM	2081	CB	GLU A	262	1	9.049		7.758	56.			29.		6
ATOM	2082	CG	GLU A	262		8.496		3.931	57.			28.		6
ATOM	2083	CD	GLU A			8.449		3.687	58.		1.00			6
ATOM	2084	OE1				8.175		7.548	59.		1.00			
ATOM	2085	OE2	GLU A			8.661		0.661						8
ATOM	2086	c	GLU A						59.3		1.00			8
ATOM	2087	Õ	GLU A 2			8.526		.754	54.8		1.00			6
	2088	N	ASP A 2			8.690		.199	53.		1.00			8
ATOM						8.778		.486	55.		1.00		-	7
MOTA	2089	CY	ASP A			9.245		.582	54.3		1.00			6
MOTA	2090	CB	ASP A 2			0.354	42	.672	54.6	541	1.00	38.4	12	6
ATOM	2091	CG	ASP A 2		20	0.982	41	.847	53.5	38	1.00	39.3	33	6
ATOM	2092		ASP A 2		22	2.064	11	.263	53.7	762	1.00	38.8	36	8
MOTA	2093	OD2	ASP A 2	63	20	3.384	:1	.779	52.4	143	1.00	39.5	60	8
MOTA	2094	С	ASP A 2	63	18	3.046		.775	53.6		1.00			6
ATOM	2095	0	ASP A 2	63		7.474		.966	54.3		1.00			8
ATOM	2096	N	TYR A 2			7.673		.002	52.3		1.00			7
ATOM	2097	CA	TYR A 2			5.508								
ATOM	2098							.357	51.7		1.00			6
			TYR A 2			.031		.149	50.5		1.00			6
MOTA		CG	TYR A 2			.824		.939	49.2		1.00			6
MOTA	2100	CDI	TYR A 2	64		.510		. 897 [.]	48.4		1.00	65.7	3	6
ATOM	2101	CEI	TYR A 2	64	17	.230	41	.709	47.2	26	1.00	68.3	5	6
ATOM			TYR A 2		17	.882	43	.786	48.9	62	1.00	66.2	3 .	['] 6
MOTA			TYR A 2		18	.611		.606	47.7	80	1.00			6
MOTA	2104		TYR A 2			.279		.570	46.9		1.00		-	6
MOTA			TYR A 2			.989		.411	45.7		1.00			8
MOTA			TYR A 2			.665		.888	51.4		1.00			6
MOTA			TYR A 2			.663		. 185					_	8
			LEU A 2				_		51.2		1.00			
MOTA						.897		.400	51.3		1.00			7
ATOM			LEU A 26			.051		.984	51.0		1.00		-	6
MOTA			LEU A 26			.474		.646	50.5		1.00		-	6
ATCM			LEU A 26			.905		.211	49.1		1.00			6
ATOM	2112 (CD1 1	LEU A 26	55	21	.176	38.	535	48.7	03	1.00	26.9	7	6
									•					

MOTA	2113	CI	2 LEU A 265	:	18.828	38.954	48.141	1.00 3	4.41	6
MOTA	2114	С	LEU A 265		17.665	38.131	52.219	1.00 3	7.41	6
MOTA	2115	0	LEU A 265		18.125		52.370	1.00 3	7.96	8
MOTA	2116		SER A 266		L6.804		53.066	1.00 3	6.21	7
MOTA	2117				16.294		54.253	1.00 3		6
ATOM -					17.263		55.427	1.00 3		6
ATOM	2119				17.190		55.991	1.00 3		8
MOTA	2120		SER A 266		14.997		54.653	1.00 3		6
MOTA	2121		SER A 266		14.889			1.00 3	_	8
MOTA	2122		LYS A 267		4.018		55.093	1.00 3		7
MOTA	2123		_		2.750		55.532	1.00 3		6
MOTA	2124 2125				1.596 1.503		55.183	1.00 3		6
MOTA	2125				.1.453		53.705-	1.00 3		6 6
ATOM ATOM	2127	CE			1.369		52.869 51.389	1.00 4		6
ATOM	2128	NZ			1.503		50.569	1.00 4		7 -
MOTA	2129	c	LYS A 267		2.791		57.043	1.00 3		6
ATOM	2130	ŏ	LYS A 267		1.758		57.694	1.00 3		8
ATOM	2131	N	PHE A 268		3.998		57.595	1.00 3		7
ATOM	2132	CA			4.192		59.016	1.00 3		6
ATOM	2133	CB	PHE A 268		5.477		59.495	1.00 3		6
ATOM	2134	CG	PHE A 268	1	5.379	36.839	59.604	1.00 34	1.54	6.
MOTA	2135	CD:	1 PHE A 268		6.506		59.940	1.00 3	5.04	6
ATOM	2136	CD:			4.161		59.429	1.00 34	1.57	6
MOTA	2137	CE:			6.423	34.691	60.108	1.00 35		6
ATOM	2138	CE			4.066	34.784	59.594	1.00 36		6
ATOM	2139	CZ	PHE A 268		5.201	34.040	59.936	1.00 34		6
MOTA	2140	C	PHE A 268		4.319	40.530	59.190	1.00 30		6
ATOM	2141 2142	0	PHE A 268 ASN A 269		4.983 3.693	41.192	58.394 60.222	1.00 30		8 7
ATOM ATOM	2142	N CA	ASN A 269		3.760	41.081 42.527	60.222	1.00 32		6
ATOM	2144	CB	ASN A 269	_	2.344	43.115	60.570	1.00 37		6
ATOM	2145	CG	ASN A 269		1.478	42.809		1.00 40		6
ATOM	2146		. ASN A 269		1.830	43.148	58.227	1.00 43		8
MOTA	2147	ND2			0.335	42.165	59.594	1.00 39		7
ATOM	2148	С	ASN A 269	1	4.553	42.854	61.710	1.00 35	.45	6
MOTA	2149	0	ASN A 269	1	4.095	43.621	62.560	1.00 41	47	8
ATOM	2150	N	LEU A 270		5.747	42.285	61.827	1.00 33		7
MOTA	2151	CA	LEU A 270		6.571	42.510	63.004	1.00 30		6
ATOM	2152	CB	LEU A 270		7.638	41.431	63.114	1.00 27		6
MOTA	2153	CG	LEU A 270		7.140	40.002	62.988	1.00 23		6
MOTA	2154		LEU A 270 LEU A 270		8.222 5.855	39.106	63.543	1.00 27		6 6
ATOM	2155 2156	CDZ	LEU A 270		7.258	39.801 43.856	63.772 63.033	1.00 28		6
ATOM ATOM	2157	ō	LEU A 270		7.347	44.554	62.017	1.00 36		8
ATOM	2158	N	SER A 271		7.749	44.207	64.216	1.00 30		7
ATOM	2159	CA	SER A 271		3.465	45.457	64.424	1.00 30		6
ATOM	2160	CB	SER A 271		7.816	46.249	65.562	1.00 29		6
MOTA	2161	OG	SER A 271	17	7.712	45.471	66.739	1.00 30	.43	8
ATOM	2162	С	SER A 271	19	9.911	45.109	64.768	1.00 33	∵09	6
ATOM	2163	0	SER A 271	20).194	43.972	65.172	1.00 29		8
ATOM	2164	N	ASN A 272		821		64.586	1.00 32		7
MOTA	2165	CA	ASN A 272		2.234	45.846	64.896	1.00 31		6
ATOM	2166	CB	ASN A 272		.036	47.141	64.771	1.00 33		6
MOTA	2167	CG	ASN A 272		.101	47.658	63.361	1.00 37		6
ATOM	2168		ASN A 272		.719	48.686	63.100	1.00 36		8
MOTA	2169		ASN A 272		460	46.952	62.437	1.00 44		7
MOTA	2170	C	ASN A 272 ASN A 272		369	45.333	66.321 66.565	1.00 32		6 8
MOTA	2171 2172	O N	VAL A 273		.970 .803	44.283 46.091	67.257	1.00 27		7
MOTA	2172	CA	VAL A 273		.839	45.741	68.668	1.00 35		6
ATOM ATOM	2174	CB	VAL A 273		.928	46.660	69.481	1.00 37		6
ATOM	2175		VAL A 273		.987	46.276	70.964	1.00 39		6
	2176		VAL A 273		.356	48.112	69.275	1.00 38		6
ATOM	2177	c	VAL A 273		.416	44.300	68.908	1.00 34		6
ATOM	2178	0	VAL A 273		.060	43.580	69.679	1.00 35	.96	8
		_	٠.				•			

ATCM	2179	N	ALA A 27	4	20.328	43.889	68.262	1.00 32.36	7
ATCM	2180				19.834				
ATOM	2181		_		18.574				6
ATOM	2182		ALA A 27						6
ATCM	2183				20.923				6
			ALA A 27		21.323				8
ATCM	2184		PHE A 27.		21.401		66.655	1.00 30.05	7
ATOM	2185				22.467	41.102	66.036	1.00 31.30	6
MOTA	2186		B PHE A 27	5	22.932	41.810	64.751	1.00 31.54	6
MOTA	2187	CG	PHE A 27	5	23.938	41.029	63.941		6
ATCM	2188	CI	1 PHE A 27	5	23.597	39.809	63.365		6
ATOM	2189	CE	2 PHE A 27	5	25.219		63.729		6
MOTA	2190	CE	1 PHE A 279	5	24.513		62.586		6
MCTA	2191	CE	2 PHE A 275	5	26.149		62.950		
ATCM	2192				25.793		62.378		6
MOTA	2193		PHE A 275		23.632				6
ATOM	2194		PHE A 27		24.252		67.040		6
ATOM	2195		LEU A 276				67.200		8
ATOM	2196				23.908		67.726		7
					24.988		68.698		6
ATOM	2197				25.221	43.594	69.141		6
ATCM	2198	CG			26.415	43.908	70.050	—	6
ATOM	2199		1 LEU A 276		26.683	45.391	70.025		6
ATOM	2200		2 LEU A 276		26.147	43.433	71.467	1.00 39.09	6
ATOM	2201	С	LEU A 276		24.682	41.244	69.894	1.00 32.79	6
MOTA	2202	0	LEU A 276		25.560	40.530	70.371	1.00 30.74	8
ATOM	2203	N	LYS A 277		23.445	41.273	70.384	1.00 33.95	7
MOTA	2204	CA	LYS A 277		23.086	40.413	71.505	1.00 36,56	6
ATOM	2205	CB	LYS A 277		21.623	40.588	71.902	1.00 35.76	6
ATOM	2206	CG	LYS A 277		21.343	41.842	72.687	1.00 42.31	6
ATOM	2207	CD	LYS A 277		20.743	41.508	74.049	1.00 45.72	6
ATOM	2208	CE	LYS A 277		21.665	40.601	74.865	1.00 47.87	6
ATOM	2209	NZ	LYS A 277		21.140	40.378	76.244	1.00 44.36	7
MCTA	2210	C	LYS A 277		23.302	38.974	71.092		
ATOM	2211	ō	LYS A 277		23.875			1.00 37.98	6
ATOM	2212	N	ALA A 278		22.832	38.179	71.845	1.00 37.54	8
ATOM	2213	CA	ALA A 278			38.654	69.886	1.00 37.17	7
ATOM	2214	CB			22.952	37.311	69.323	1.00 34.51	6
			ALA A 278		22.638	37.341	67.820	1.00 35.38	6
ATOM	2215	C	ALA A 278		24.368	36.831	69.550	1.00 30.63	6
ATOM	2216	0	ALA A 278		24.605	35.790	70.167	1.00 27.62	8
ATOM	2217	N	PHE A 279		25.303	37.624	69.049	1.00 29.24	7
ATCM	2218	CA	PHE A 279		26.722	37.347	69.167	1.00 31.48	6
ATOM	2219	CB	PHE A 279		27.490	38.558	68.645	1.00 33.25	6
ATOM	2220	CG	PHE A 279		28.974	38.396	68.663	1.00 39.28	6
ATCM	2221		PHE A 279		29.578	37.337	68.000	1.00 41.15	6
ATOM	2222	CD2	-		29.776	39.328	69.315	1.00 40.66	6
ATOM	2223	CE1			30.960	37.209	67.987	1.00 44.22	6
ATOM	2224	CE2	PHE A 279		31.153	39.213	69.378	1.00 41.38	6
ATOM	2225	cz	PHE A 279		31.750	38.152	68.614	1.00 44.52	6
ATCM	2226	С	PHE A 279		27.116	37.043	70.6:1	1.00 31.81	6
ATOM	2227	0	PHE A 279		27.627	35.953	70.935	1.00 27.51	8
MCTA	2228	N	ASN A 280		26.860	38.005	71.503	1.00 29.32	7
ATOM	2229	CA	ASN A 280		27.192	37.851	72.907	1.00 29.32	6
MCTA	2230	CB	ASN A 280		26.927	39.153			
ATOM	2231	CG	ASN A 280				73.660	1.00 30.39	6
					27.907	40.245	73.278	1.00 30.68	6
ATOM	2232		ASN A 280		29.117	40.030	73.303	1.00 33.34	8
MOTA	2233		ASN A 280		27.395	41.419	72.931	1.00 27.00	7
ATOM	2234	C	ASN A 280		26.524	36.680	73.616	1.00 30.01	6
MCTA	2235	0	ASN A 280		27.167	36.004	74.419	1.00 29.58	8
ATOM	2236	N	ILE A 281		25.252	36.423	73.335	1.00 30.46	7
MCTA	2237	CA	ILE A 281		24.594	35.291	73.983	1.00 33.71	6
ATCM	2238	CB	ILE A 281		23.107	35.161	73.569	1.00 36.14	6
MCIA	2239	CG2	ILE A 281		22.541	33.820	74.032	1.00 36.18	6
ATCM	2240		ILE A 281		22.298	36.307	74.177	1.00 33.52	6
ATCM	2241		ILE A 281		20.835	36.243	73.834	1.00 37.16	6
MOT	2242	c	ILE A 281		25.330	34.006	73.631	1.00 34.06	6
ATOM	2243	ŏ	ILE A 281		25.385	33.071	74.437	1.00 31.94	8
NOM	2244	N	VAL A 282		25.896	33.960			7
	~~ 14	**	TALL A LUL		40.000	JJ. JOU	72.427	1.00 35.31	,

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					•								
ATOM	224	5 C	A .VAL	A 282		26.6	54	32.785	72.	005	1.00	36.45	6
MOTA				A 282		27.0		32.871				35.62	
MOTA			G1 VAL			27.8		31.604		126	1.00	31.20	
MOTA			G2 VAL			25.8		33.080				34.51	-
MOTA	224: 225:			A 282 A 282		27.9		32.723	72.1			37.80	_
MOTA MOTA	225			A 283		28.1		31.722	73.5			36.12	_
ATOM	225			A 283		28.6 29.9		33.808 33.884	72.8 73.9			38.45	
ATOM	225			A 283		30.5		35.272	73.4			40.06 39.38	-
ATOM	225			A 283		30.9		35.625	72.0			41.90	6 6
ATOM	225	5 C1	D ARG	A 283		31.4		37.048	71.9			41.36	6
MOTA	2256			A 283		32.6	47	37.206	72.8			43.35	7
MOTA	2257			A 283		33.1	62	38.373	73.2	215		42.55	6
ATOM	2258		II ARG			32.62		39.516	72.7			39.95	7
MOTA	2259 2260		12 ARG	A 283 A 283		34.22		38.392	74.0			41.72	7
MOTA MOTA	2261			A 283 A 283		29.63		33.587	75.0			40.01	6
ATOM	2262			A 284		30.35 28.50		32.862 34.141	75.7			39.01	8
MOTA	2263			A 284		28.08		33.923	75.5 76.8			40.30	7 6
ATOM	2264			A 284		26.75		34.647	77.1			47.53	6
ATOM	2265	CG	GLU.	A 284		26.87		36.176	77.0			56.10	6
MOTA	2266		_	A 284	:	25.54	12	36.923	77.1			60.77	6
ATOM	2267		1 GLU			24.65		36.682	76.3	29		61.41	8
MOTA	2268			A 284		25.38		37.763	78.0	96	1.00	62.21	8
MOTA	2269 2270			A 284		27.95		32.429	77.1			40.72	6
ATOM ATOM	2270			A 284 A 285		28.56		31.922	78.1			45.29	8
ATOM	2272	CA		A 285		27.18 26.97		31.721 30.288	76.3 76.5			34.82	7
ATOM	2273	CB		A 285		25.84		29.752	75.6			30.84	6 6
MOTA	2274	CG				5.69		28.253	75.8			22.95	6
MOTA	2275	CG	2 VAL A	A 285	2	4.54	5	30.433	75.9			26.26	6
MOTA	227.6	С	VAL A		2	8.18	1	29.366	76.3	41		31.93	6
ATOM	2277	0	VAL A			8.49		28.556	77.2	14	1.00	33.46	8
ATOM	2278 2279	N	PHE A			8.84		29:466	75.1			29.43	7
ATOM ATOM	2279	CA CB	PHE A			9.97		28.586	74.9			24.26	6
ATOM	2281	CG	PHE A			9.83 8.60		27.957 27.095	73.5			22.57	6
ATOM	2282		l PHE A			7.40		27.639	72.8			23.46	6 6
ATOM	2283	CD:				8.66		25.718	73.6			21.95	6
ATOM	. 2284		PHE A	286	2	6.28	1	26.814	72.6			24.90	6
MOTA	2285	CE				7.54		24.892	73.4	11	1.00	18.06	6
MOTA	2286	CZ	PHE A			6.35		25.437	72.9			20.23	6
ATOM	2287 2288	C O	PHE A			1.36		29.200	74.99			25.14	6
MOTA MOTA	2289	Ŋ	GLY A			2.330 1.480		28.566	74.50			23.16	8
ATOM	229.	CA	GLY A			2.783		30.416 31.065	75.52 75.63			25.51 26.86	7 6
ATOM	2291	С	GLY A			3.353		31.511-	74.27			26.28	6
MOTA	2292	0	GLY A			2.644		31.549	73.27	_		26.29	8
MOTA	2293	N.	GLU A			4.637		31.849	74.23			27.17	7
MOTA	2294	CA	GLU A			5.274		32.291	72.99		1.00	33.20	6
MOTA	2295	CB	GLU A			6.680		32.828	73.26	59	1.00	35.09	6
ATOM	2296	CG.				5.726		34.104	74.08		1.00		6
ATOM ATOM	2297 2298	CD OF1	GLU A GLU A			5.970		35.231	73.42		1.00		6
ATOM	2299		GLU A			5.221 5.130		35.493	72.22		1.00		8
ATOM	2300	c	GLU A			5.386	_	35.858 31.204	74.09 71.93		1.00		8 6
MOTA	2301	Ō	GLU A			5.596		30.029	72.24		1.00		8
ATOM	2302	N	GLY A			.268		31.619	70.66		1.00		7
MOTA	2303	CA	GLY A			.373		30.698	69.54		1.00		6
MOTA	2304	С	GLY A			.948	3	31.372	68.30		1.00		5
MOTA	2305	0	GLY A			. 556		32.437	68.39		1.00		8
MOTA	2306	N	VAL A			.764		30.758	67.14		1.00	_	7
MOTA	2307	CA	VAL A			.277		31.349	65.90		1.00		6
MOTA	2308 2309	CB CG1	VAL A VAL A			616		30.301	65.03		1.00 2		6
MOTA MOTA	2310		VAL A			.616 .100		30.976	63.81 65.85		1.00 2 1.00 1		6 6
ATOM			· · · · · ·		20	. 100	•	29.632		4	1.00	33	J

•		_	VAL A	200	35.137	31.975	65.105	1.00 25.97	6
ATOM	2311	C				31.279	64.672	1.00 22.32	8
ATOM	2312	0	VAL A		34.218				
MOTA	2313	N	TYR A	291	35.217	33.293	64.914	1.00 27.33	7
ATOM	2314	CA	TYR A	291	34.188	34.052	64.203	1.00 26.69	6
	2315	CB	TYR A		33.925	35.356	64.939	1.00 25.51	6
MOTA						35.178	66.435	1.00 28.73	6
MOTA	2316	CG	TYR A		33.935			1.00 29.72	6
ATOM	2317	CD1	TYR A	291	35.025	35.596	67.191		
MOTA	2318	CE1	TYR A	291 [']	35.059	35.414	68.563	1.00 29.53	6
	2319		TYR A		32.874	34.565	67:094	1.00 27.39	6
ATOM		CE2			32.898	34.377	68.466	1.00 31.03	6
MOTA	2320					34.808	69.194	1.00 31.85	6
MOTA	2321	CZ	TYR A		33.997			1.00 38.03	8
MOTA	2322	OH	TYR A		34.030	34.647	70.562		
MOTA	2323	С	TYR A	291	34.527	34.345	62.745	1.00 27.99	6
ATOM	2324	٠٥	TYR A	291	35.608	34.843	62.415	1.00 24.83	8
	2325	Ŋ	LEU A		33.567	34.042	61.880	1.00 30.17	7
MOTA					33.726	34.220	60.441	1.00 28.26	6
MOTA	2326	CA	LEU A				59.741	1.00 27.70	6
ATOM	2327	CB	LEU A		33.561	32.861			6
ATOM	2328	CG	LEU A	292	34.191	31.643	60.435	1.00 24.64	
MOTA	2329	CD1	LEU A	292	33.867	30.380	59.661	1.00 24.66	6
	2330	CD2	LEU A	292	35.686	31.825	60.553	1.00 23.19	6
MOTA	2331	C	LEU A		32.649	35.175	59.944	1.00 25.59	6
MOTA					31.640	35.394	60.611	1.00 18.11	8
MOTA	2332	0	LEU A				58.770	1.00 28.55	7
ATOM	2333	N	GLY A		32.869	35.749			6
ATOM	2334	CA	GLY A	293	31.878	36.653	58.223	1.00 31.26	
ATOM'	2335	С	GLY A	293	30.722	35.815	57.714	1.00 34.84	6
ATOM	2336	0	GLY A	293	30.463	34.724	58.234	1.00 34.11	8
	2337	Ň	GLY A		30.036	36.312	56.689	1.00 35.34	7
ATOM		CA	GLY A		28.918	35.581	56.124	1.00 34.84	6
ATOM	2338		GLY A		28.142	36.445	55.155	1.00 34.79	6
MOTA	2339	C				37.473	54.699	1.00 37.05	8
MOTA	2340	0	GLY A		28.644				7
ATOM	2341	N	GLY A		26.917	36.035	54.842	1.00 31.87	
MOTA	2342	CA	GLY A	295	26.102	36.806	53.925	1.00 27.78	6
ATOM	2343	С	GLY A	295	25.969	38.245	54.378	1.00 27.09	6
	2344	.0	GLY A	295	26.192	38.558	55.546	1.00 27.03	8
ATOM	2345	N	GLY A		25.596	39.119	53.450	1.00 24.67	7
ATOM			GLY A		25.440	40.527	53.757	1.00 25.28	6
MOTA	2346	CA				41.262	52.446	1.00 27.64	6
ATOM	2347	С	GLY A		25.562			1.00 26.65	8
MOTA	2348	0	GLY A		26.591	41.163	51.771		7
ATOM	2349	N	TYR A	297	24.526	42.009	52.078	1.00 30.21	
MOTA	2350	CA	TYR A	297	24.543	42.704	50.801	1.00 30.62	6
ATOM	2351	CB	TYR A	297	23.560	42.011	49.859	1.00 29.50	6
	2352	CG	TYR A		23.717	40.516	49.953	1.00 30.33	6
MOTA			TYR A		23.174	39.810	51.031	1.00 30.86	6
MOTA	2353				23.450	38.449	51.226	1.00 30.74	6
MOTA	2354	CE1	TYR A				49.062	1.00 31.20	6
ATOM	2355	CD2	TYR A		24.538	39.824		1.00 32 08	6
ATOM	2356	CE2			24.821	38.460	49.247		
MOTA	2357	cz	TYR A	297	24.275	37.781	50.332	1.00 30.92	6
ATOM	2358	OH	TYR A	297 .	24.539	36.440	50.509	1.00 29 60	8
	2359	C	TYR A	297	24.267	44.195	50.875	1.00 32.07	6
MOTA	2360	ō	TYR A		24.134	44.849	49.840	1.00 33.83	8
ATOM					24.180	44.725	52.094	1.00 31.41	7
ATOM	2361	N	HIS A	290			52.289	1.00 33.94	6
MOTA	2362	CA	HIS A	298	23.961	46.153		1.00 34.75	6
ATOM	2363	CB	HIS A	298	22.761	46.430	53.194		
MOTA	2364	CG	HIS A	298	22.379	47.880	53.256	1.00 35.16	6
	2365	CD2	HIS A	298	22.558	48.809	54.224	1.00 35.72	6
ATOM		NO1	HIS A	298	21.779	48.538	52.205	1.00 34.10	7
MOTA	2366	CE1	HIS A	298	21.605	49.809	52.522	1.00 31.84	6
MOTA	2367	(E1	TID A	200			53.742	1.00 35.46	7
ATOM	2368		HIS A	270 200	22.069	50.000	52.962	1.00 36.21	6
ATOM	2369	С	HIS A	298	25.213	46.697			
ATOM	2370	0	HIS A		25.471	46.405	54.133	1.00 33.83	8
	2371	N	PRO A		25.992	47.519	52.234	1.00 36.69	7
ATOM	2372	CD	PRO A	299	25.680	47.997	50.881	1.00 35.57	6
MOTA			PRO A		27.238	48.142	52,689	1.00 35.17	6
λ TOM	2373	CA	PRO A	200	27.586	49.073	51.525	1.00 37.75	6
ATOM	2374	CB					50.954	1.00 37.76	6
MOTA	2375	CG	PRO A		26.216	49.399		1.00 34.47	6
MOTA	2376	С	PRO A	299	27.045	48.886	54.000	1.00 34.4/	•

MOTA	237	7 0	PRO	O A 299		27.78	1 48.670	54.963	1.00 33.67	
MOTA				R A 300		26.05				8
			_							
ATOM				R A 300		25.74				6
MOTA	238	0 C	B TYP	R A 300		24.49	6 51.377	7 55.009	1.00 35.56	6
ATOM	238	1 C	G TYF	R A 300		24.64	8 52.524	54.028		6
ATOM-				R A 300		25.37				
										6
MOTA	238			R A 300		25.46				6
MOTA	238			R A 300		24.01	6 53.752	54.259	1.00 35.92	6
MOTA	238	5 C	E2 TYF	00E A 3		24.09	8 54.793	53.334	1.00 36.78	6
ATOM	238			A 300		24.82				
										6
MOTA	238			A 300		24.92				8
MOTA	238	8 C		A 300		25.49	7 49.546	56.3.69	1.00 31.10	6
ATOM	238:	90	TYR	A 300	-	26.063	49.692	57.440	1.00 30.62	8
ATOM	239	и о		A 301		24.66				7
ATOM	239			A 301		24.32				
										6
ATOM	239			A 301		23.21		56.624	1.00 24.69	6
ATOM	239	3 C	ALA	A 301		25.539	46.727	57.552	1.00 30.06	6
ATOM	239	4 0	ALA	A 301		25.848	3 46.579			8
MOTA				A 302		26.223		56.557		
										7
MOTA	2396			A 302		27.404			1.00 30.55	6
MOTA	2397		3 LEU	A 302		28.012	45:002	55.441	1.00 31.83	6
ATOM	2398	3 CG	LEU	A 302		29.315	44.223	55.323	1.00 30.01	6
MOTA	2399	CE	1 LEU	A 302		29.491			1.00 32.09	6
ATOM	2400			A 302		30.475				
									1.00 32.23	6
ATOM	2401			A 302		28.418		5,7.663	1.00 29.79	6
ATOM	2402	0	LEU	A 302		28.796	45.676	58.746	1.00 27.68	8
ATOM	2403	N	ALA	A 303		28.842	47.299	57.179	1.00 27.92	7
ATOM	2404	CA	A.IA	A 303		29.818		57.877	1.00 25.00	
ATOM	2405			A 303						6
						30.026		57.137	1.00 23.62	6
MOTA	2406			A 303		29.397	48.397	59.305	1.00 25.06	6
ATOM	2407	0	ALA	A 303		30.088	48.015	60.248	1.00 26.90	8
MOTA	2408	N	ARG	A 304		28.258	49.054	59.472	1.00 24.06	7
ATOM	2409			A 304		27.794		60.810	1.00 24.37	6
	2410			A 304 .						0
ATOM						26.420	50.052	60.758	1.00 23.99	6
MOTA	2411			A 304		26.328	51.257	59.815	1.00 28.77	6
ATOM	2412	CD	ARG	A 304		25.106	52.089	60.156	1.00 29.96	6
ATOM	2413	NE	ARG	A 304		23.943	51.233	60.369	1.00 36.43	7
ATOM	2414	CZ		A 304		22.893	51.573	61.110	1.00 37.01	6
	2415	NH:		A 304						
ATOM						22.854	52.757	61.713	1.00 36.37	7
ATOM	2416	NH:		A 304		21.896	50.719	61.269	1.00 34.36	7
MOTA	2417	С	ARG .	A 304		27.727	48.142	61.691	1.00 24.24	6
MOTA	2418	0	ARG	A 304		28.343	48.099	62.762	1.00 22.34	8
ATOM	2419	N		A 305		26.994	47.132	61.221	1.00 24.51	7
	2420			A 305						
ATOM		CA				26.801	45.883	61.959	1.00 22.70	6
ATOM	2421	CB		A 305		25.880	44.960	61.175	1.00 18.13	6
MOTA	2422	С	ALA .	. 305		28.089	45.142	62.351	1.00 23.33	. 6
ATOM	2423	0	ALA .	. 305		28.237	44.725	63.506	1.00 21.51	8
MOTA	2424	N		. 306		29.016	44.961	61.411	1.00 22.79	7
	2425							01.411		
ATOM		CA		A 306		30.244	44.270	61.764	1.00 24.33	6
MOTA	2426	CB		A 306		31.029	43.842	60.524	1.00 26.93	6
ATOM	2427	ÇG	TRP A	A 306		30.604	42.503	59.952	1.00 27.96	6
ATOM	2428	CD2	TRP A	306		30.861	42.013	58.629	1.00 26.38	6
	2429		TRP A			30.366				_
MOTA							40.688	58.570	1.00 24.96	6.
MOTA	2430		TRP A			31.462	42.563	57.490	1.00 23.00	6
ATOM	2431		TRP A			29.983	41.484	60.620	1.00 28.53	6
MOTA	2432	NE1	TRP A	306		29.837	40.392	59.797	1.00 25.62	7
ATOM	2433		TRP A			30.450	39.904	57.414	1.00 24.51	
										6
MOTA	2434	CZ3	TRP A			31.548	41.784	56.343	1.00 25.37	6
MOTA	2435	CH2	TRP A			31.042	40.465	56.315	1.00 24.20	6
ATOM	2436	С	TRP A	306		31.129	45.108	62.676	1.00 26.51	6
MOTA	2437	ō	TRP A			31.908	44.570	63.464	1.00 25.07	8
	2438		THR A							
ATOM		N				31.003	46.427	62.575	1.00 28.08	7
MOTA	2439	CA	THR A			31.785	47.323	63.415	1.00 27.91	6.
ATOM ·	2440	CB	THR A	. 307		31.484	48.796	63.100	1.00 27.86	6
ATOM	2441	OG1		307		31.994	49.119	61.799	1.00 30.17	8
ATOM	2442	CG2				32.120			1.00 24.72	6
	244	C G Z	THY Y			JE. 12U	49.704	64.137	1.00 24.72	J
		•		•						

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Figure 17-38

	ATOM	2443	C	THR A 30	7	31.441	47.041	64.863	1.00 29.35	6
	ATOM	2444	ō	THR A 30		32.316	46.989	65.725	1.00 32.56	8
	MOTA	2445	Ŋ	LEU A 30		30.159	46.857	65.135	1.00 30.60	7
	MOTA	2446	CA			29.740	46.555	66.490	1.00 33.69	6
	ATOM	2447	CB	LEU A 30		28.256	46.215	66.525	1.00 33.69	
	MOTA	2448	CG	LEU A 30		27.338	47.337	66.058		6
		2449		1 LEU A 30		25.903			1.00 34.79	6
	MOTA			2 LEU A 30			46.887	66.153	1.00 31.98	6
	MOTA	2450				27.569	48.542	66.924	1.00 36.96	6
	MOTA	2451	C	LEU A 308		30.531	45.353	66.965	1.00 34.57	6
	MOTA	2452	0	LEU A 308		31.230	45.417	67.975	1.00 33.75	8
	ATOM	2453	N	ILE A 309		30.423	44.262	66.208	1.00 35.78	7
	ATOM	2454	CA	ILE A 309		31.108	43.017	66.540	1.00 35.87	6
	MOTA	2455	CB	ILE A 309		30.939	41.949	65.431	1.00 34.95	6
	MOTA	2456	CG:			31.733	40.695	65.799	1.00 31.21	6
	MOTA	2457	CG:			29.445	41.631	65.212	1.00 34.25	6
	ATOM	2458		1 ILE A 309		28.726	41.014	66.407		6
	MOTA	2459	C	ILE A 309		32.589	43.238	66.772	1.00 35.81	6
	MOTA	2460	0	ILE A 309		33.183	42.617	67.657	1.00 38.19	8
	MOTA	2461	N	TRP A 310		33.197	44.111	65.977	1.00 36.10	7
	MOTA	2462	CA	TRP A 310		34.612	44.384	66.169	1.00 35.26	6
	MOTA	2463	CB	TRP A 310	1	35.150	45.311	65.075	1.00 32.61	6
	MOTA	2464	CG	TRP A 310		36.619	45.588	65.220	1.00 30.79	6
	ATOM	2465	CD2	2 TRP A 310		37.679	44.620	65.274	1.00 29.93	6
	MOTA	2466	CE2	2 TRP A 310		38.882	45.330	65.474	1.00 28.42	6
	ATOM	2467	CE3	3 TRP A 310		37.731	43.224	65.174	1.00 31.59	6
	ATOM	2468	CD1	L TRP A 310		37.206	46.804	65.380	1.00 30.62	6
	ATOM	2469	NE1	TRP A 310		38.565	46.659	65.536	1.00 29.37	7
	ATOM	2470	CZ2	TRP A 310		40.126	44.691	65.578	1.00 27.91	6
	ATOM	2471	CZ3	TRP A 310		38.978	42.585	65.279	1.00 28.06	6
	ATOM-	2472	CH2	TRP A 310		40.150	43.322	65.479	1.00 26.50	6
	ATOM	2473	С	TRP A 310		34.744	45.040	67.545	1.00 36.00	6
	MOTA	2474	0	TRP A 310		35.365	44.476	68.440	1.00 36.24	8
	ATOM	2475	N	CYS A 311		34.134	46.213	67.715	1.00 34.57	7
	ATOM	2476	CA	CYS A 311	•	34.183	46.937	68.985	1.00 32.82	6
	ATOM	2477	ĊВ	CYS A 311		33.169	48.085	68.996	1.00 35.62	6
	ATOM	2478	SG	CYS A 311		33.439	49.401	67.796	1.00 32.36	16
	ATOM	2479	C	CYS A 311		33.912	46.061	70.206	1.00 32.01	6
	MOTA	2480	0	CYS A 311		34.452	46.313	71.280	1.00 29.82	8
	ATOM	2481	N	GLU A 312		33.062	45.049	70.053	1.00 32.57	7
	MOTA	2482	CA	GLU A 312		32.731	44.159	71.171	1.00 33.86	6
	ATOM	2483	CB	GLU A 312		31.557	43.252	70.807	1.00 34.19	6
	ATOM	2484	CG	GLU A 312		30.442	43.185	71.844	1.00 40.27	6
	ATOM	2485	CD	GLU A 312		30.923	42.821	73.239	1.00 43.80	6
	ATOM	2486	OE1			31.685	41.831	73.383	1.00 44.81	8
	MOTA	2487	OE2			30.516	43.522	74.195	1.00 41.54	8
	ATOM	2488	c	GLU A 312		33.953	43.298	71.456	1.00 33.77	6
	ATOM	2489	ō	GLU A 312		34.253	42.957	72.603	1.00 32.07	8
	ATOM	2490	N	LEU A 313		34.647	42.945	70.382	1.00 33.45	7
	ATOM	2491	CA	LEU A 313		35.848	42.135	70.473	1.00 32.89	6
	ATOM	2492	CB	LEU A 313		36.172	41.513	69.115	1.00 32.14	6
	ATOM	2493	CG	LEU A 313		35.154	40.493	68.626	1.00 27.73	6
		~ 4 ~ 4		LEU A 313		35.587	39.956	67.269	1.00 30.39	6
	ATOM	2494		LEU A 313		35.053	39.367	69.648	1.00 27.87	6
	MOTA	2496	C	LEU A 313		36.976	43.031		1.00 31.64	6
	MOTA			LEU A 313		37.605	42.793	71.925	1.00 31.74	8
	MOTA	2497	O N						1.00 31.74	
	ATCM	2498		SER A 314		37.206	44.064	70.099		7
	MOTA	2499	CA	SER A 314		38.232	45.067	70.328	1.00 35.59 1.00 36.47	6
	MOTA	2500	CB	SER A 314		38.107	46.154	69.256		6
	MOTA	2501	OG C	SER A 314		39.141	47.120	69.353	1.00 44.55	8
	ATCM	2502	C	SER A 314		38.046	45.661	71.730	1.30 37.82	6
	MOTA	2503	0	SER A 314		39.015	46.043	72.395	1.00 37.32	8
	ATCM	2504	N	GLY A 315		36.794	45.725	72.175	1.00 38.05	7
_	ATOM	2505	CA	GLY A 315		36.506	46.243	73.498	1.00 42.42	6
	MOTA	2506	C	GLY A 315		36.295	47.744	73.568	1.00 46.80	6
	ATOM	2507	0	GLY A 315		35.923	48.276	74.618	1.00 47.85	8
	ATCM	2508	N	ARG A 316		36.518	48.438	72.458	1.00 48.90	7

MOTA	2509	CA ARG A 316		36.340	49.88	5 72.448	3 1.00 52.27	6
MOTA	2510	CB ARG A 3:16		37.14				6
ATOM	2511	CG ARG A 316		36.730				
	2512							6
ATOM				37.734			1.00 53,76	6
ATOM	2513	NE ARG A 316		39.028	3 49.854	4 69.019	1.00 53.67	7
ATOM	- 2514	CZ ARG A 316		40.135	50.223	68.383		6
MOTA	2515	NH1 ARG A 316		40.110				7
ATOM	2516	NH2 ARG A 316		41.266				
ATOM	2517	C ARG A 316						7
				34.882				6
MOTA	2518	O ARG A 316		34.075	49.781	71.652	1.00 54.96	8
MOTA	2519	N GLU A 317		34.547	51.361	73.182	1.00 51.87	7
MOTA	2520	CA GLU A.317		33.185	51.900			6
MOTA	2521	CB GLU A 317	-	33.111				
ATOM	2522	CG GLU A 317		32.549				6
	2523							6
ATOM		CD GLU A 317		33.353				6
ATOM	2524	OE1 GLU A 317		33.025	51.741	77.556	1.00 64.59	8
ATOM	2525	OE2 GLU A 317		34.305	51.302	75.832	1.00 68.64	8
MOTA	2526	C GLU A 317		32.642			1.00 51.27	6
ATOM	2527	O GLU A 317		33.270				
ATOM		N VAL A 318					1.00 49.34	8
				31.457			1.00 51.30	7
MOTA		CA VAL A 318		30.780	51.962	70.280	1.00 48.80	6
MOTA		CB VAL A 318		29.522	51.071	70.169	1.00 47.11	6
ATOM	2531	CG1 VAL A 318		28.875	51.237	68.808	1.00 45.53	6
ATOM	2532	CG2 VAL A 318		29.895	49.631		1.00 47.05	6
ATOM		C VAL A 318		30.349	53.411			
ATOM		VAL A 318					1.00 47.64	6
				29.511	53.867		1.00 47.61	8
MOTA		N PRO A 319		30.925	54.165	69.234	1.00 48.14	7
MOTA		D PRO A 319		31.960	53.836	68.247	1.00 48.87	6
ATOM	2537 (CA PRO A 319		30.538	55.569	69.093	1.00 52.54	6
MOTA	2538 (CB PRO A 319		31.438	56.051	67.954	1.00 49.96	6
ATOM	2539	G PRO A 319	•	31.612	54.802	67.141		
ATOM		PRO A 319					1.00 50.17	6
		DBO 3 310		29.052	55.679	68.764	1.00 55.84	6
MOTA	2541 0			28.531	54.913	67.953	1.00 56.06	8
MOTA		I GLU A 320		28.369	56.624	69.402	1.00 59.20	7
MOTA	2543 C	A GLU A 320		26.942	56.804	69.167	1.00 62.61	6
MOTA	2544 0	B GLU A 320		26.302	57.588	70.313	1.00 65.59	6
ATOM	2545 C	G GLU A 320		26.727	59.042	70.365	1.00 73.01	6
ATOM		D GLU A 320		26.007				
		E1 GLU A 320			59.823	71.451	1.00 76.93	6
ATOM				24.755	59.832	71.446	1.00 77.37	8
ATOM		E2 GLU A 320		26.697	60.431	72.303	1.00 79.46	8
ATOM	2549 C			26.698	57.551	67.863	1.00 61.40	6
ATOM	2550 O	GLU A 320		25.663	58.197	67.699	1.00 62.33	8
ATOM	2551 N	LYS A 321		27.650	57.463	66.939	1.00 59.47	7
ATOM	2552 C			27.519	58.150			
ATOM	2553 C			27.340		65.662	1.00 59.54	6
					59.648	65.897	1.00 61.36	. 6
MOTA	2554 C			23.620	60.323	66.366	1.00 65.23	6
ATOM	2555 C	_		13.169	59.691	67.643	1.00 66.59	6
ATOM	2556 C	Ē LYS A 321		E).564	60.215	67.960	1.00 67.34	6
MOTA	2557 N	Z LYS A 321		30.591	61.699	68.100	1.00 68.58	7
ATOM	2558 C	LYS A 321		28.766	57.941			
	2559 0					64.806	1.00 59.24	6
ATOM		LYS A 321		29.845	57.623	65.319	1.00 58.70	8
ATOM	2560 N			28.608	58.146	63.500	1.00 57.55	7
ATOM	2561 C	A LEU A 322		29.702	58.002	62.543	1.00 54.72	6
ATOM	2562 CI			29.171	57.450	61.214	1.00 52.96	6
ATOM	2563 CC			28.141	56.316	61.295	1.00 52.97	
		01 LEU A 322			50.310			6
ATOM				27.708	55.932	59.899	1.00 49.01	6
MOTA		2 LEU A 322		28.716	55.115	62.035	1.00 54.73	6
MOTA	2566 C	LEU A 322		30.250	59.406	62.313	1.00 53.47	-6
ATOM	2567 O	LEU A 322		29.512	60.383	62.464	1.00 53.39	8
ATOM	2568 N	ASN A 323		31.530	59.521	61.965	1.00 51.43	7
	2569 CA		٠.					
ATOM			٠	32.089	60.842	61.706	1.00 50.32	6
ATOM	2570 CB			33.591	60.905	62.035	1.00 52.31	6
ATOM	2571 CG			34.428	59.964	61.189	1.00 55.06	6
MOTA	2572 OD	1 ASN A 323		34.386	58.744	61.363	1.00 55.77	8
ATOM		2 ASN A 323		35.195	60.530	60.259	1.00 52.71	7
ATOM	2574 C	ASN A 323		31.843	61.199	60.243	1.00 48.63	6
ATOM	-3,4	-1017 11 343		-I.043	01.133	-00.243	7.00 40.03	o
		•					•	

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а пом	2575	0	ASN A 323	31.135	60.479	59.538	1.00 47.52	8
ATOM								
ATOM	2576	N	ASN A 324	32.426		59.792	1.00 47.66	7
ATOM	257 7	CA	asn a 324	32.242	62.769	58.419	1.00 49.25	6
ATOM	2578	CB	ASN A 324	32.758	64.200	58.292	1.00 50.73	6
	2579	CG	ASN A 324	32.025		59.205	1.00 53.83	6
MOTA								
ATOM	2580	OD1	. ASN A 324	30.812		59.096	1.00 56.90	8
ATOM	2581	ND2	ASN A 324	32.755	65.789	60.119	1.00 54.93	7
ATOM	2582	C	ASN A 324	32.906	61.891	57.367	1.00 49.34	6
				32.275		56.379	1.00 47.22	8
ATOM	2583	0	ASN A 324					
ATOM	2584	N	LYS A 325	34.182	61.590	57.586	1.00 48.27	7
ATOM	2585	CA	LYS A 325	34.957	60.759	56.676	1.00 46:57	6
ATOM	2586	CB	LYS A 325	36.314	60.453	57.305	1.00 49.94	6
			LYS A 325	37.299		56.399	1.00 54.75	6
MOTA	2587	CG			-			
ATOM	2588	CD	LYS A 325	38.562		57.173	1.00 58.22	6
ATOM	2589	CE	LYS A 325	39.236	60.521	57.844	1.00 58.35	6
ATOM	2590	NZ	LYS A 325	40.473	60.128	58.566	1.00 59.81	7
	2591	c	LYS A 325	34.202		56.410	1.00 44.85	6
ATOM								
MOTA	2592	0	LYS A 325	34.065		55.263	1.00 43.59	. 8
MOTA	2593	N	ALA A 326	33.712	58.843	57.483	1.00 42.27	7
ATOM	2594	CA	ALA A 326	32.964	57.597	57.387	1.00 40.91	6
	2595	CB	ALA A 326	32.663	57.067	58.773	1.00 36.86	6
ATOM								6
MOTA	2596	С	ALA A 326	31.666		56.612	1.00 43.30	
ATOM	2597	0	ALA A 326	31.342	57.028	55.705	1.00 42.83	8
ATOM	2598	N	LYS A 327	30.918	58.843	56.977	1.00 45.56	7
ATOM	2599	CA	LYS A 327	29.657	59.146	56.306	1.00 47.23	б
				29.023	60.407	56.892	1.00 49.59	6
ATOM	2600	CB	LYS A 327					
ATOM	2601	CG	LYS A 327	28.547		58.329	1.00 54.63	6
ATOM	2602	CD	LYS A 327	· 28.024	61.591	58.862	1.00 55.89	6
ATOM	2603	CE	LYS A 327	27.529	61.483	60.299	1.00 58.28	6
	2604	NZ	LYS A 327	26.304		60.426	1.00 59.91	7
MOTA								6
ATOM	2605	С	LYS A 327	29.888	59.347	54.816	1.00 46.97	
MOTA	2606	0	LYS A 327	29.090	58.913	53.990	1.00 48.10	8
ATOM	2607	N	GLU A 328	30.986	60.012	54.480	1.00 44.99	7
ATOM	2608	CA	GLU A 328	31.325	60.264	53.091	1.00 43.18	6
			GLU A 328	32.417		53.027	1.00 47.93	6
MOTA	.2609	CB						
MOTA	2610	CG	GLU A 328	31.993	62.621	53.710	1.00 53.65	6
ATOM	2611	CD	GLU A 328	33.112	63.630	53.831	1.00 55.79	6
ATOM	2612	OE1	GLU A 328	33.642	64.060	52.783	1.00 58.73	8
	2613	OE2	GLU A 328	33.459	63.991	54.979	1.00 58.13	8
ATOM				31.789	58.971	52.437	1.00 41.56	6
ATOM	2614	C	GLU A 328					
ATOM	2615	0	GLU A 328	31.537	58.743	51.255	1.00 39.41	8
ATOM	2616	N	LEU A 329	32.465	58.123	53.211	1.00 40.64	7
ATOM	2617	CA	LEU A 329	32.940	56.844	52.695	1.00 36.45	6
	2618	CB	LEU A 329	33.623	56.032	53.801	1.00 34.70	6
ATOM						53.433	1.00 35.69	Ğ
ATOM	2619	CG	LEU A 329	34.100	54.610			
ATOM	2620	CD1	LEU A 329	35.195	54.642	52.359	1.00 30.11	6
ATOM	2621	CD2	LEU A 329	34.619	53.926	54.683	1.00 34.63	6
ATOM	2622	С	LEU A 329	31.746	56.064	52.157	1.00 35.77	6
		ŏ	LEU A 329	31.692	55.746	50.975	1.00 34.94	8
ATOM	2623						1.00 34.78	7
MOTA	2624	Ν.		30.784	55.770	53.029		
MOTA	2625	CA	LEU A 330	29.599	55.028	52.630	1.00 34.95	6
ATOM	2626	CB	LEU A 330	28.631	54.914	53.803	1.00 30.95	6
	2627	CG	LEU A 330	29.164	54.115	54.991	1.00 32.66	6
ATOM				28.051	53.904	56.022	1.00 31.74	6
ATOM	2628	CDI	LEU A 330					
MOTA	2629	CD2	LEU A 330	29.674	52.769	54.509	1.00 30.73	6
MOTA	2630	С	LEU A 330	28.877	55.631	51.428	1.00 37.28	6
ATOM	2631	0	LEU A 330	28.395	54.901	50.557	1.00 40.56	8
			LYS A 331	28.806	56.957	51.383	1.00 38.24	7
MOTA	2632	N						6
ATOM	2633	CA	LYS A 331	28.140	57.661	50.294	1.00 39.59	9
ATOM	2634	CB	LYS A 331	27.994	59.146	50.643	1.00 42.31	ε
ATOM	2635	CG	LYS A 331	27.129	59.399	51.873	1.00 45.93	6
			LYS A 331	27.017	60.879	52.244	1.00 49.72	6
ATOM	2636	CD	TIO W 33T				1.00 53.66	6
MOTA	2637	CE	LYS A 331	26.271	61.698	51.193		9
ATOM	2638	NZ	LYS A 331	26.053	63.114	51.640	1.00 54.22	7
ATOM	2639	С	LYS A 331	28.863	57.514	48.958	1.00 41.02	6
		ō	LYS A 331	28.220	57.485	47.904	1.00 39.58	8
atom	2640	9	TTO WALL	20.240	37.403			_

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Figure 17-41

-				30 400	E	40 005		-
MOTA	2641	N	SER A 332	30.192	57.413	49.005	1.00 42.69	7
MOTA	2642	CA	SER A 332	30.998	57.274	47.792	1.00 46.02	6
	2643	CB	SER A 332	32.494	57.243	48.124	1.00 47.51	6
MOTA			SER A 332	32.862	56.066	48.823	1.00 50.97	8
MOTA	2644	OG						6
MOTA	2645	С	SER A 332	30.634	56.007	47.040	1.00 48.51	
MOTA	2646	0	SER A 332	30.706	55.959	45.811	1.00 49.45	8
MOTA	2647	N	ILE A 333	30.241	54.982	47.786	1.00 51.56	7
ATOM	2648	CA	ILE A 333	29.869	53.713	47.187	1.00 54.86	6
		CB	ILE A-333	29.657	52.626	48.246	1.00 55.80	6
MOTA	2649			29.388	51.285	47.559	1.00 52.34	6
MOTA	2650	CG2						
MOTA	2651		ILE A 333	30.892	52.522	49.140	1.00 56.89	6
MOTA	2652	CD1	ILE A 333	30.766	51.456	50.204	1.00 60.31	6
ATOM	2653	C	ILE A 333	28.579	53.813	46.396	1.00 57.07	6
MOTA	2654	Ó	ILE A 333	27.572	54.321	46.897	1.00 55.59	8
	2655	N	ASP A 334	28.623	53.320	45.160	1.00 61.14	7
MOTA				27.456	53.300	44.281	1.00 65.55	6
MOTA	2656	CA	ASP A 334					
MOTA	2657	CB	ASP A 334	27.888	53.259	42.811	1.00 69.19	6
MOTA	2658	CG	ASP A 334	28.784	52.073	42.491	1.00 70.21	6
ATOM	2659	OD1	ASP A 334	29.097	51.875	41.298	1.00 71.47	8
ATOM	2660		ASP A 334	29.181	51.344	43.427	1.00 70.91	8
	2661	C	ASP A 334	26.660	52.041	44.627	1.00 65.65	6
MOTA				26.797	50.996	43.990	1.00 63.91	8
MOTA	2662	0	ASP A 334				•	7
ATOM	2663	N	PHE A 335	25.822	52.153	45.649	1.00 65.73	
ATOM	2664	CA	PHE A 335	25.041	51.021	46.104	1.00 63.44	6
ATOM	2665	CB	PHE A 335	24.980	51.034	47.632	1.00 58.05	6
MOTA	2666	CG	PHE A 335	24.039	50.028	48.195	1.00 53.82	6
	2667		PHE A 335	24.178	48.679	47.886	1.00 52.40	6
ATOM			PHE A 335	22.978	50.429	48.989	1.00 51.33	6
ATOM	2668			23.265	47.742	48.356	1.00 52.73	6
MOTA	2669	CE1						
MOTA	2670	CE2		22.062	49.503	49.462	1.00 53.20	6
ATOM	2671	CZ	PHE A 335	22.204	48.151	49.144	1.00 51.76	6
MOTA	2672	С	PHE A 335	23.629	50.893	45.535	1.00 65.55	6
ATOM	2673	0	PHE A 335	23.230	49.810	45.097	1.00 67.33	8
	2674	N	GLU A 336	22.874	51.986	45.537	1.00 66.47	7
ATOM			GLU A 336	21.497	51.948	45.048	1.00 67.43	6
MOTA	2675	CA				43.626	1.00 71.79	6
ATOM	2676	CB	GLU A 336	21.422	51.379			
MOTA	2677	CG	GLU A 336	19.982	51.245	43.116	1.00 78.77	6
MOTA	2678	CD	GLU A 336	19.868	50.505	41.789	1.00 82.67	6
ATOM	2679	OE1	GLU A 336	20.232	49.306	41.734	1.00 83.29	8
ATOM	2680		GLU A 336	19.410	51.126	40.801	1.00 84.26	8
	2681	C	GLU A 336	20.655	51.069	45.971	1.00 64.72	6
ATOM				20.686	49.840	45.876	1.00 59.84	8
ATOM	2682	0	GLU A 336				1.00 64.47	7
MOTA	2683	N	GLU A 337	19.901	51.710	46.858		
ATOM	2684	CA	GLU A 337	19.045	51.003	47.805	1.00 65.83	6
ATOM	2685	CB	GLU A 337	18.398	52.003	48.759	1.00 64.20	6
ATOM	2686	CG	GLU A 337	17.753	51.370	49.964	1.00 64.26	6
ATOM	2687	CD	GLU A 337	18.774	50.€JO	50.850	-1.00 64.04	6
	2688	OE1		19.741	51.3.2	51.261	1.00 61.66	8
MOTA	2689		GLU A 337	18.608	49.483	51.132	1.00 63.64	8
ATOM		OE2				47.063	1.00 67.13	6
ATOM	2690	C	GLU A 337	17.950	50.239			8
MOTA	2691	0	GLU A 337	17.269	50.807	46.205	1.00 68.27	
ATOM	2692	N	PHE A 338	17.779	48.960	47.394	1.00 67.22	7
ATOM	2693	CA	PHE A 338	16.764	48.129	46.748	1.00 68.05	6
	2694	CB	PHE A 338	16.445	46.919	47.626	1.00 69.68	6
ATOM			PHE A 338	15.228	46.158	47.187	1.00 72.35	6
ATOM	2695	CG					1.00 72.37	6
ATOM	2696		PHE A 338	15.122	45.674	45.888	1.00 74.37	
MOTA	2697		PHE A 338	14.172	45.941	48.074	1.00 73.61	6
ATOM	2698	CE1	PHE A 338	13.980	44.984	45.478	1.00 73.39	6
ATOM	2699		PHE A 338	13.024	45.250	47.672	1.00 73.26	6
	2700	CZ	PHE A 338 -	12.929	44.771	46.369	1.00 73.34	6
ATOM			PHE A 338	15.481	48.902	46.434	1.00 68.45	6
ATOM	2701	C	FUE W 330			_	1.00 67.92	8
ATOM	2702	0	PHE A 338	. 15.286	49.367	45.310		7
. ATOM	2703	N	ASP A 339	14.606	49.026	47.426	1.00 68.98	
ATOM	2704	CA	ASP A 339	13.358	49.759	47.261	1.00 70.68	6
ATOM	2705	CB	ASP A 339	12.596	49.758	48.588	1.00 71.06	6
	2706	CG	ASP A 339	11.381	50.678	48.581	1.00 72.05	6
ATOM	~,00	<u> </u>	227					

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									_
ATOM	2707	OD1	. ASP A 339		11.548	51.893	48.320	1.00 72.29	8
ATOM	2708	OD2	ASP A 339		10.262	50.188	48.858	1.00 71.03	8
	2709	C	ASP A 339		13.715	51.183	46.853	1.00 73.18	6
MOTA			ASP A 339			51.884	47.592	1.00 73.78	8
MOTA	2710	0			14.407				
ATOM	2711	Ŋ	ASP A 340		13.247	51.600	45.677	1.00 76.36	7
ATOM	2712	CA	ASP A 340		13.518	52.943	45.152	1.00 78.34	6
	2713	CB	ASP A 340		12.410	53.385	44.189	1.00 77.55	6
ATOM			-	,			-	1.00 78.90	6
\mathtt{MOTA}	2714	CG	ASP A 340		12.462	52.655	42.864		
MOTA	2715	OD1	ASP A 340		12.348	51.408	42.855	1.00 78.38	8
ATOM	2716	OD2	ASP A 340		12.620	53.336	41.830	1.00 78.74	8
		C	ASP A 340		13.687	54.017	46.214	1.00 79.51	6
MOTA	2717			•				1.00 80.19	8
MOTA	2718	0	ASP A 340		14.587	54.856	46.117	-	
ATOM	2719	N	GLU A 341		12.824	54.000	47.224	1.00 79.64	7
ATOM	2720	CA	GLU A 341		12.922	54.998	48.271	1.00 80.05	6
	2721	СВ	GLU A 341		12.269	56.301	47.811	1.00 83.75	6
MOTA						57.442	48.806	1.00 89.02	6
MOTA	2722	CG	GLU A 341		12.411				
ATOM	2723	· CD	GLU A 341		11.756	58.724	48.328	1.00 91.52	6
ATOM	2724	OE1	GLU A 341		10.515	58.738	48.175	1.00 93.33	. 8
	2725	OE2			12.484	59.716	48.102	1.00 92.65	8
MOTA						54.578	49.597	1.00 77.98	6
ATOM	2726	С	GLU A 341		12.317			_	
MOTA	2727	0	GLU A 341		11.102	54.610	49.777	1.00 79.82	8
ATOM	2728	N	VAL A 342		13.179	54.181	50.523	1.00 74.49	7
	2729	CA	VAL A 342		12.745	53.793	51.859	1.00 71.55	6
ATOM			VAL A 342		13.224	52.383	52.245	1.00 72.40	6
MOTA	2730	CB							
MOTA	2731	CG1			12.672	52.004	53.610	1.00 71.16	6
ATOM	2732	CG2	VAL A 342		12.797	51.391	51.207	1.00 74.35	6
ATOM	2733	С	VAL A 342		13.454	54.778	52.766	1.00 68.46	6
	2734	ŏ	VAL A 342		12.952	55.154	53.829	1.00 68.96	8
ATOM						55.184	52.311	1.00 61.61	7
ATOM	2735	N	ASP A 343		14.636				
ATOM	2736	CA	ASP A 343		15.486	56.114	53.029	1.00 54.91	6
MOTA	2737	CB	ASP A 343		14.678	57.303	53.543	1.00 55.06	6
	2738	CG	ASP A 343		15.556	58.390	54.114	1.00 54.44	6
ATOM					15.002	59.351	54.694	1.00 56,20	8
ATOM	2739	OD1							8
MOTA	2740	QD2		•	16.795	58.287	53.969	1.00 49.32	
ATOM	2741	С	ASP A 343		16.152	55.401	54.198	1.00 50.85	6
MOTA	2742	0	ASP A 343		15.557	55.209	55.257	1.00 49.32	8
	2743	N	ARG A 344		17.396	55.004	53.980	1.00 47.84	7
MOTA					18.195	54.321	54.981	1.00 45.34	6
MOTA	2744	CA	ARG A 344						
ATOM	2745	CB	ARG A 344		18.883	53.099	54.358	1.00 45.00	6
ATOM	2746	CG	ARG A 344		17.950	51.969	53.974	1.00 38.03	6
	2747	CD	ARG A 344		17.185	51.531	55.188	1.00 35.83	6
ATOM		NE	ARG A 344		16.278	50.439	54.885	1.00 39.20	7
MOTA	2748							1.00 39.89	6
ATOM	2749	CZ	ARG A 344		15.350	49.993	55.724		
ATOM	2750	NH1	ARG A 344		15.217	50.561	56.917	1.00 40.17	7
ATOM	2751	NH2	ARG A 344		14.566	48.976	55.375	1.00 40.75	7
	2752	С	ARG A 344		19.250	55.278	55.515	1.00 44.72	6
ATC I			ARG A 344		20.170	54.869	56.223	1.00 46.97	8
ATC:M	2753	0						1.00 45.81	7
ATOH	2754	N	SER A 345		19.113	56.552	55.157		
ATOM	2755	CA	SER A 345		20.045	57.596	55.577	1.00 43.66	6
ATOM	2756	CB	SER A 345		19.538	58.960	55.115	1.00 43.44	6
	2757	OG	SER A 345		18.292	59.260	55.722	1.00 45.62	8
ATOM								1.00 42.79	6
ATOM	2758		SER A 345		20.258	57.627	57.089		
ATOM	2759	O	SER A 345		21.364		57.552	1.00 42.62	8
ATOM	2760	N	TYR A 346		19.200	57.354 ·	57.851	1.00 40.55	7
	2761	CA	TYR A 346		19.280	57.352	59.308	1.00 41.05	6
ATOM						56.819	59.905	1.00 41.74	6
ATOM	2762	CB	TYR A 346		17.971				
ATOM	2763	CG	TYR A 346		17.668	55.355	59.630	1.00 43.47	6
ATOM	2764	CD1	TYR A 346		18.331	54.333	60.328	1.00 44.45	6
	2765	CE1	TYR A 346		18.044	52.983	60.088	1.00 41.02	6
ATOM					16.710	54.988	58.682	1.00 42.08	6
ATOM	2766	CD2	TYR A 346						
ATCM	2767	CE2	TYR A 346		16.416	53.644	58.434	1.00 40.59	6
ATCM	2768	CZ	TYR A 346		17.086	52.649	59.139	1.00 41.66	5
	2769	OH	TYR A 346		16.806	51.324	58.884	1.00 39.60	8
ATCM			TYR A 346		20.466	.56.517	59.796	1.00 42.92	6
ATCM	2770	C					60.799	1.00 42.65	8
ATCM	2771	0	TYR A 346		21.101	56.844			
ATCM	2772	N	MET A 347		20.757	55.443	59.067	1.00 44.59	7
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7. MOM	2773	ÇA	MET 2	A 347		21.859	54.546	59.388	1 00	45.89	6
MOTA											
ATOM	2774	CB	WEI I	A 347		21.950	53.433	58.353	1.00	45.51	6
ATOM -	. 2775	CG	MET	A 347		20.727	52.567	58.244	1.00	45.01	6
MOTA	2776	SD	MET A	A 347		21.062	51.258	57.066	1.00	47.13	16
ATOM	2777	CE	MET A	A 347		21.545	52.237	55.676	1 00	44.55	6
ATOM -	2778	C		347		23.188	55.286	59.404		48.47	6
MOTA	2779	0	MET A	347		24.129	54.888	60.098	1.00	49.49	8
	_										7
MOTA	2780	N		A 348		23.259	56.351	58.610		49.03	
MOTA	2781	CA	LEU A	348		24.458	57.178	58.499	1.00	48.86	6
-	2782	CB		348		24.355	58.082	57.269		45.51	6
MOTA											
ATOM	2783	CG	LEU A	348		24.280	57.424	55.895	1.00	44.47	6
	2784	CD	1 LEU A	3/18		23.908	58.476	54.859	1 00	43.62	6
MOTA											
MOTA	2785	CD:	2 LEU A	348	-	25.618	56.757	55.565	1.00	42.53	6
ATOM	2786	С	LEU A	348		24.644	58.049	59.738	1.00	49.33	6
						-					
ATOM	2787	ο.				25.765	58.369	60.123		49.78	8
MOTA	2788	N	GLU A	349		23.537	58.428	60.358	1.00	48.34	7.
	2789	CA	GLU A			23.591	59.279	61.533		49.24	6
MOTA											
MOTA	2790	CB	GLU A	349		22.198	59.848	61.811	1.00	48.36	6
	2791	CG	GLU A	349		21.628	60.584	60.607	7 00	45.52	6
MOTA											
ATOM	2792	CD	GLU A			22.598	61.619	60.065		42.94	6
ATOM	2793	OE:	L GLU A	349		22.934	62.560	60.812	1.00	40.82	8
MOTA	2794		GLU A			23.028	61.483	58.900		38.30	8
ATOM	2795	С	GLU A	349		24.119	58.531	62.745	1.00	48.32	6
	2796	0	GLU A	349		25.226	58.783	63.219	1 00	47.87	8
ATOM											
MOTA	2797	N	THR A	350		23.325	57.602	63.248	1.00	48.97	7
ATOM	2798	CA	THR A	350		23.744	56.832	64.398	1.00	50.70	6
								65.342			ě
ATOM .	2799	CB	THR A			22.558	56.596			51.02	
MOTA	2800	OG]	. THR A	350		22.071	57.865	65.803	1.00	49.11	8
	2801	CG2	THR A	350		22.983	55.763	66.537	1 00	51.58	6
MOTA											
MOTA	2802	C	THR A			24.361	55.507	63.954		49.56	6
MOTA	2803	0	THR A	350	•	23.979	54.947	62.923	1.00	50.55	8
	2804	N	LEU A			25.333	55.028	64.725	1 00	46.88	7
MOTA											
ATOM	2805	CA	LEU A	351		26.018	53.781	64.417	1.00	45.35	6
ATOM	2806	CB	LEU A	351		27.342	53.726	65.185	1.00	47.05	6
MOTA	2807	CG	LEU A			28.257	52.502	65.072		49.54	6
ATOM	2808	CD1	LEU A	351		29.575	52.777	65.766	1.00	51.50	6
	2809	CD2	LEU A	351		27.603	51.302	65.692	1 00	48.35	6
MOTA											
MOTA	2810	С	LEU A	351		25.145	52.584	64.772	1.00	44.79	6
MOTA	2811	0	LEU A	351		25.131	51.578	64.061	1.00	41.45	8
			LYS A			24.420	52.711	65.880		45.27	7
ATOM .	2812	N									
ATOM	2813	CA	LYS A	352		23.531	51.662	66.375	1.00	44.62	6
	2814	CB	LYS A	352		23.764	51.464	67.873	1.00	42.23	6
MOTA	_										
MOTA	2815	CG	LYS A			25.197	51.075	68.187		44.94	6
ATOM	2816	CD	LYS A	352		25.572	51.262	69.650	1.00	46.80	6
	2817	CE	LYS A	352		24.765	50.389	70.581	1 00	45.79	6
MOTA											
ATOM	2818	NZ	LYS A	352		25.236	50.586	71.975	1.00	47.31	7
ATOM	2819	С	LYS A	352		22.096	52.087	66.116	1.00	45.12	6
							53.236	65.756		47.07	8
MOTK	2820	0	LYS A			21.837					
ATOM	2821	N	ASP A	353		21.162	51.161	66.285	1.00	44.62	7 ·
	2822	CA	ASP A	353		19.761	51.474	66.060	1 00	46.43	6
MOTA											
MOTA	2823	CB	ASP A	353		19.302	50.943	64.692	1.00	49.38	6
MOTA	.2824.	CG	ASP A	353		19.813	49.546	64.396	.1.00	51.52	6
											8
MOTA	2825	OD1				21.028	49.396	64.158	1.00		
ATOM	2826	OD2	ASP A	353		19.005	48.596	64.398	1.00	52.35	8
	2827	С	ASP A	353		18.841	50.968	67.165	1.00	45.90	6
MOTA											
ATOM	2828	0	ASP A	353	-	19.152	50.001	67.854	1.00		8
	2829	N	PRO A	354	•	17.687	51.629	67.348	1.00	45.86	7
ATOM								66.587	1.00		
MOTA	2830	CD	PRO A			17.162	52.775				6
MOTA	2831	CA	PRO A	354	1	16.723	51.243	68.378	1.00	45.52	6
	2832	СВ	PRO A			15.585	52.245	68.159	1.00		6
MOTA											~
MOTA	2833	CG	PRO A		1	15.681	52.513	66.664	1.00		6
	2834	С	PRO A	354	1	16.277	49.804	68.188	1.00	44.13	6
ATOM			PRO A			16.352		67.078	1.00		8
ATOM	2835	0					49.271			_	
ATCM ·	2836	N	TRP A	355	3	15.821	49.174	69.267	1.00	42.77	7
	2837	CA	TRP A		1	15.358	47.801	69.168	1.00	43.35	6
ATOM						14.982		70.539	1.00		6
ATOM	2838	CB	TRP A	ررر		14.502	47.225	, 0.333	2.00		•

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ATOM	283	9 C	G TRP A 355	16.168	46.752	71.322	1.00 52.43	_
MOTA	284		D2 TRP A 355	16.574				6
ATOM	284		E2 TRP A 355	17.789				6
ATOM	284							6
			E3 TRP A 355	16.031				6
ATOM	2843		D1 TRP A 355	17.125			1.00 54.39	6
MOTA	2844		El TRP A 355	18.103	46.731	72.468	1.00 57.31	7
ATOM	2845	5 C:	Z2 TRP A 355	18.469	44.249	72.602		6
ATOM	2846	5 C:	Z3 TRP A 355	16.706				6
MOTA	2847	7 CI	12 TRP A 355	17.913				
ATOM	2848		TRP A 355	14.177				6
MOTA	2849		TRP A 355	13.508				6
ATOM	2850							8
			ARG A 356	13.942				7
ATOM	2851			12.855				6
ATOM	2852			13.413	46.044	65.451	1.00 35.06	6
ATOM	2853			14.120	47.308	64.976	1.00 32.47	6
MOTA	2854	CI	ARG A 356	14.969	47.082	63733	1.00 29.54	6
ATOM	2855	NE	ARG A 356	15.600	48.323	63.296	1.00 28.91	7
ATOM	2856	CZ	ARG A 356	16.514	48.403	62.335	1.00 30.60	6
ATOM	2857	NH		16.916	47.305	61.702	1.00 33.52	
ATOM	2858			17.020	49.582	61.996		7
ATOM	2859		ARG A 356	12.270	44.879		1.00 30.10	7
ATOM	2860		ARG A 356			67.361	1.00 36.01	6
				12.447	43.831	66.742	1.00 38.38	8
MOTA	2861		GLY A 357	11.587	44.949	68.499	1.00 36.04	7
MOTA	2862	CA		11.001	43.758	69.085	1.00 36.08	6
MOTA	2863	С	GLY A 357	9.514	43.596	68.851	1.00 34.51	6
MOTA	2864	0	GLY A 357	8.943	44.196	67.943	1.00 36.77	8
MOTA	2865	N	GLY A 358	8.892	42.772	69.687	1.00 36.04	7
ATOM	2866	CA	GLY A 358	7.466	42.506	69.593	1.00 32.26	6
MOTA	2867	С	GLY A 358	7.106	41.263	70.385	1.00 29.85	6
ATOM	2868	0	GLY A 358	7.832	40.839	71.288	1.00 28.86	8
MOTA	2869	N	GLU A 359	5.975	40.667	70.055	1.00 30.88	7
ATOM	2870	CA	GLU A 359	5.550	39.455	70.743	1.00 32.58	6
ATOM	2871	CB	GLU A 359	4.034	39.289	70.604	1.00 38.60	
ATOM	2872	CG	GLU A 359	3.230	40.435	71.222		6
ATOM	2873	CD	GLU A 359	1.957	40.762		1.00 47.44	6
ATOM	2874	OE:		1.123		70.445	1.00 50.93	6
ATOM	2875	OE:			39.852	70.221	1.00 52.13	8
	2876	C C		1.798	41.942	70.061	1.00 51.03	8
ATOM	2877		GLU A 359	6.250	38.275	70.091	1.00 28.29	6
ATOM		0	GLU A 359	6.790	38.382	68.997	1.00 27.88	8
MOTA	2878	N	VAL A 360	6.263	37.147	70.772	1.00 27.97	7
ATOM	2879	CA	VAL A 360	6.859	35.957	70.193	1.00 25.86	6
ATOM	2880	CB	VAL A 360	7.673	35.168	71.237	1.00 22.02	6
ATOM	2881	CG1		8.155	33.849	70.641	1.00 19.45	б
ATOM	2882	CG2		8.850	36.009	71.698	1.00 17.88	6
ATOM	2883	С	VAL A 360	5.703	35.099	69.670	1.00 28.04	6
ATOM	2884	C	VAL A 360	4.842	34.655	70.440	1.00 27.34	8
ATOM	2885	N	ARG A 361	5.663	34.898	68.358	1.00 27.70	7
ATOM -	2886	C.,	ARG A 361	4.612	34.091	67.765	1.00 32.85	6
MOTA	2887	CB	ARG A 361	4.693	34.164	66.242	1.00 32.30	6
MOTA	2888	CG	ARG A 361	4.243	35.504	65.687	1.00 38.81	6
ATOM	2889	CD	ARG A 361	4.546	35.653	64.201	1.00 40.09	6
ATOM	2890	NE	ARG A 361	5.974	35.812	63.961		
	2891	cz	ARG A 361				1.00 38.77	7
MOTA			ARG A 361	6.514	35.989	62.763	1.00 39.21	6
MOTA	2892			5.748	36.027	61.685	1.00 40.79	7
ATOM	2893		ARG A 361	7.822	36.145	62.643	1.00 42.94	7
ATOM	2894	С	ARG A 361	4.689	32.638	68.222	1.00 35.65	6
MOTA	2895	0	ARG A 361	5.768	32.097	68.471	1.00 37.08	8
ATOM	2896	N	LYS A 362	3.526	32.017	68.347	1.00 37.80	7
MOTA	2897	CA	LYS A 362	3.436	30.626	68.757	1.00 39.91	6
ATOM	2898	CB	LYS A 362	1.982	30.152	68.648	1.00 43.43	6
ATOM	2899	CG	LYS A 362	1.014	30.803	69.640	1.00 45.11	6
ATOM	2900	CD	LYS A 362	1.117	32.346	69.673	1.00 49.43	<i>-</i>
ATOM	2901	CE	LYS A 362	0.813	33.022	68.327	1.00 45.02	6 6
ATOM	2902	NZ	LYS A 362	0.962	34.509	68.422	1.00 45.02	0
ATOM ATOM	2903	C	LYS A 362	4.320	29.809	67.831		7
	2904	Ö	LYS A 362	4.953			1.00 39.25	6
MOTA	2702	•		4.733	28.835	68.248	1.00 35.45	8

		-				
MOTA	1 2905 N GLU A 363	4.35	8 30.229	9 66.56	8 1.00 41.13	
ATOM	1 2906 CA GLU A 363	5.14			9 1.00 43.96	
ATOM		5.22		64.27		
ATOM		3.89				
MOTA		4.04				
ATOM		4.57	1 31.224			
MOTA		3.64				8
ATOM		6.55		66.046	5 1.00 42.39	ě
ATOM		7.06		65.989	1.00 41.48	ē
MOTA		7.18			0 1.00 37.27	7
ATOM		8.53			1.00 35.30	6
ATOM		9.03			1.00,36,88	6
MOTA		10.44				6
ATOM ATOM		9.01				6
ATOM		8.650				6
ATOM		9.622				8
ATOM	2922 CA LYS A 365	7.664		69.165		7
ATOM	2923 CB LYS A 365	7.674 6.598				6
MOTA	2924 CG LYS A 365	6.826		71.358		6
ATOM	2925 CD LYS A 365	5.837		71.899		6
ATOM	2926 CE LYS A 365	6.120		72.995 73.509		6
MOTA	2927 NZ LYS A 365	5.191		74.585		6
ATOM	2928 C LYS A 365	7.452		70.007		7
MOTA	2929 O LYS A 365	8.195		70.442		6
MOTA	2930 N ASP A 366	6.427		69.209	1.00 29.85	8 7
MOTA	2931 CA ASP A 366	6.115		68.807		6
ATOM	2932 CB ASP A 366	4.948	25.522	67.818	1.00 35.98	6
ATOM	2933 CG ASP A 366	3.711	26.206	68.381	1.00 39.88	6
ATOM	2934 OD1 ASP A 366	3.124		69.359	1.00 40.16	8
ATOM	2935 OD2 ASP A 366 2936 C ASP A 366	3.326		67.848	1.00 43.21	8
ATOM ATOM		7.343	24.866	68.161	1.00 33.07	6
MOTA		7.753	23.763	68.540	1.00 32.64	8
ATOM	2938 N THR A 367 2939 CA THR A 367	7.932	25.565	67.193	1.00 31.89	7
ATOM	2940 CB THR A 367	9.088	25.045	66.490	1.00 31.17	6
ATOM	2941 OG1 THR A 367	9.712 8.707	26.070 26.620	65.572	1.00 31.55	6
ATOM	2942 CG2 THR A 367	10.780	25.404	64.714 64.723	1.00 34.37	8
MOTA	2943 C THR A 367	10.146	24.633	67.472	1.00 33.55 1.00 33.09	6
MOTA	2944 O THR A 367	10.586	23.476	67.485	1.00 33.09	6 8
MOTA	2945 N LEU A 368	10.570	25.579	68.298	1.00 31.85	7
MOTA	2946 CA LEU A 368	11.582	25.264	69.288	1.00 32.87	6
MOTA	2947 CB LEU A 368	11.848	26.478	70.179	1.00 27.73	6
MOTA	2948 CG LEU A 368	12.887	27.449	69.588	1.00 29.05	6
MOTA	2949 CD1 LEU A 368	14.260	26.777	69.541	1.00 23.05	6
MOTA	2950 CD2 LEU A 368	12.473	27.896	68.193	1.00 26.53	6
MOTA	2951 C LEU A 368	11.157	24.053	70.107	1.00 35.16-	6
ATOM ATOM	2952 O LEU A 368 2953 N GLU A 369	11.910	23.077	70.217	1.00 35.18	8
ATOM	2953 N GLU A 369 2954 CA GLU A 369	9.942	24.101	70.649	1.00 37.56	7
ATOM	2955 CB GLU A 369	9.431 7.956	22.993	71.442	1.00 40.23	6
ATOM	2956 CG GLU A 369	7.722	23.216 24.460	71.770		6
ATOM	2957 CD GLU A 369	6.281	24.460	72.617	1.00 48.51	6
ATOM	2958 OE1 GLU A 369	5.777	23.724	73.067 73.782	1.00 51.93	6
ATOM	2959 OE2 GLU A 369	5.652	25.636	72.710	1.00 52.84 1.00 58.33	8
ATOM	2960 C GLU A 369	9.633	21.672	70.701	1.00 38.33	8
ATOM	2961 O GLU A 369	10.087	20.684	71.286	1.00 41.14	6
ATOM	2962 N LYS A 370	9.309	21.653	69.411	1.00 39.65	8 7
MOTA	2963 CA LYS A 370	9.497	20.443	68.636	1.00 38.26	6
MOTA	2964 CB LYS A 370	9.144	20.654	67.166	1.00 40.63	6
MOTA	2965 CG LYS A 370	7.675	20.597	66.854	1.00 44.49	6
MOTA	2966 CD LYS A 370	7.495	20.358	65.363	1.00 49.95	6
MOTA	2967 CE LYS A 370	6.052	20.023	65.015	1.00 54.28	6
MOTA	2968 NZ LYS A 370	5.890		63.574	1.00 55.44	7
MOTA	2969 C LYS A 370	10.948	20.034	68.730	1.00 37.85	6
MOT	2970 O LYS A 370	11.261	18.930	69.156	1.00 37.95	8

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MOTA	2971	N ALA A	371	11.827	20.944	68.325	1.00 37.78	7
ATOM	2972	CA ALA A	371	13.264	20.704	68.340	1.00 36.39	6
				14.007	22.030	68.200	1.00 37.73	6
ATOM	2973			_				
MOTA	2974	C ALA A		13.719	19.972	69.603	1.00 35.42	6
ATOM	2975	O ALA A	371	14.424	18.964	69.525	1.00 32.06	8
MOTA	2976	N ALA A	372	13.317	20.478	70.766	1.00 33.96	7
					19.848	72.024	1.00 32.22	6
MOTA	2977	CA ALAA		13.695				
MOTA	2978	CB ALA A	372	12.946	20.486	73.165	1.00 28.27	6
MOTA	2979	C ALA A	.372	13.372	18.362	71.953	1.00 31.75	6
	2980	O ALA A		14.183	17.517	72.338	1.00 31.56	8
ATOM						71.432	1.00 32.72	7 .
MOTA	2981	N ALA A		- 12.187	18.059			
ATOM	2982	CA ALA A	373	11.710	16.684	71.305	1.00 32.32	6
MOTA	2983	CB ALA A	373	10.206	16.689	71.103	1.00 30.18	6
	2984	C ALA A		12.385	15.921	70.172	1.00 33.13	6
ATOM						70.468	1.00 35.87	8
MOTA	2985	OT1 ALA A		13.078	14.926			
MOTA	2986	OT2 ALA A	373	12.218	16.320	69.003	1.00 34.11	8
ATOM	2987	ZN ZN Z	951	22.693	34.497	53.990	1.00 36.45	6
	2988	OH2 WAT S	1	35.654	44.211	49.416	1.00 9.27	8
MOTA							1.00 21.27	. 8
ATOM	2989	OH2 WAT S	2	24.480	33.130	53.069		
ATOM	2990	OH2 WAT S	3	22.124	30.277	59.314	1.00 14.69	8
ATOM	2991	OH2 WAT S	4	13.839	20.611	75.741	1.00 27.94	8
	2992	OH2 WAT S	5	34.033	41.903	46.522	1.00 44.54	8
MOTA						55.781	1.00 23.79	8
MOTA	2993	OH2 WAT S	6	15.039	42.130			
ATOM	2994	OH2 WAT S	7	32.737	41.397	75.900	1.00 15.80	8
ATOM	2995	OH2 WAT S	8	11.367	22.606	58.814	1.00 23.37	8
ATOM	2996	OH2 WAT S	9	13.909	18.160	65.105	1.00 29.93	8
				29.655	56.108	58.029	1.00 50.54	8
ATOM	2997	OH2 WAT S	10					
ATOM	2998	OH2 WAT S	11	45.405	17.964	51.885	1.00 9.28	8
ATOM	2999	OH2 WAT S	12	21.870	35.873	34.515	1.00 32.78	8
ATOM	3000	OH2 WAT S	13	43.504	35.670	33.779	1.00 28.85	8
		OH2 WAT S	14	2.054	37.997	68.430	1.00 40.53	8
ATOM	3001						1.00 21.42	.8
ATOM	3002	OH2 WAT S	15	49.730	28.024	55.966		
ATOM	3003	OH2 WAT S	16	47.503	32.289	34.336	1.00 26.13	8
ATOM	3004	OH2 WAT S	17	6.101	26.102	64.434	1.00 21.69	8
	3005	OH2 WAT S	18	10.761	46.748	45.836	1.00 15.79	8
ATOM				9.146	16.861	61.441	1.00 16.68	8
ATOM	3006	OH2 WAT S	19					
ATOM	3007	OH2 WAT S	20	5.684	34.080	76.599	1.00 37.53	8
MOTA	3008	OH2 WAT S	21	14.896	33.163	49.117	1.00 34.17	8
ATOM	3009	OH2 WAT S	22	43.346	40.839	36.825	1.00 35.64	8
			23	0.516	27.705	69.174	1.00 21.02	8
MOTA	3010					29.717	1.00 29.80	8
ATOM	3011	OH2 WAT S	24	41.270	25.444			
MOTA	3012	OH2 WAT S	25	17.818	29.142	54.584	1.00 27.92	8
ATOM	3013	OH2 WAT S	26	21.512	60.572	56.912	1.00 16.77	8
	3014	OH2 WAT S	27	21.211	33.582	48.347	1.00 23.93	8
ATOM			28	47.805	24.638	56.619	1.00 23.73	8
ATOM	3015	OH2 WAT S						8
MOTA	3016	OH2 WAT S	2.5	44.624	50.302	58.154	1.00 16.79	
MOTA	3017	OH2 WAT S	3 i`	31.096	16.437	51.311	1.00 26.61	8
ATOM	3018	OH2 WAT S	31	39.837	38.833	55.145	1.00 32.28	8
			.32	11.660	43.601	63.704	1.00 22.94	8
MOTA	3019					53.058	1.00 26.85	8
ATOM	3020	OH2 WAT S	33	49.899	23.474			
ATOM	3021	OH2 WAT S	34	34.624	17.734	32.228	1.00 21.18	8
	3022	OH2 WAT S	35	26.926	15.913	62.444	1.00 27.01	8
ATOM			36	8.893	28.686	63.905	1.00 27.68	8
MOTA	3023	OH2 WAT S			26.634	43.532	1.00 24.42	8
ATOM	3024	OH2 WAT S	37	23.381				
ATOM	3025	OH2 WAT S	38	48.484	27.990	65.270	1.00 34.86	8
	3026	OH2 WAT S	39	43.382	28,410	74.379	1.00 25.68	8
MOTA		OH2 WAT S	40	42.904	18.967	70.272	1.00 29.45	8
MOTA	302.7			20.521	53.828	50.298	1.00 22.35	8
MOTA	3028	OH2 WAT S	41					
ATOM	3029	OH2 WAT S	42	13.310	38.921	48.404	1.00 23.32	8
	3030	OH2 WAT S	43	9.787	46.265	60.012	1.00 33.51	8
ATOM		OH2 WAT S	44	36.089	30.416	51.377	1.00 47.75	8
ATOM	3031			14.831		42.151	1.00 50.96	8
ATOM	3032	OH2 WAT S	45		48.131			
ATOM	3033	OH2 WAT S	46	54.162	48.194	60.971	1.00 22.66	8
ATOM	3034	OH2 WAT S	47	38.943	61.290	63.509	1.00 33.73	8
	3035	OH2 WAT S	48	29.980	18.112	33.130	1.00 35.80	8
ATOM		OHE WAT B		31.879	50.673	44.528	1.00 24.39	8
ATOM	3036	OH2 WAT S	49	. JI,075	50.075	520	_,	~

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Figure 17-47

ATOM	303	7 OH2	WAT	S 50		39.86	3 14.62	9 64.307	1.00 24.19	8
ATOM	303		WAT			26.11	_			_
ATOM	303	_	WAT	S 52		48.07				_
MOTA	304								1.00 36.38	8
				S 53		50.80				
ATOM	304		WAT	S 54		49.54				- 8
MOTA		_	WAT			6.88	7 23.42	64.961	1.00 17.49	8
ATOM	304:	3 OH2	WAT	S 56		25.69	8 39.89	1 37.674		
ATOM	304	4 OH2	WAT	S 57		45.49	8 44.10	1 55.393		
MOTA	3049	5 OH2	WAT			44.66				_
ATOM	3046		WAT			21.91				8
	3047		WAT						1.00 26.96	8
ATOM						27.29				8
MOTA	3048		WAT		_	19.80				8
MOTA	3049		TAW		-	30.84	3 18.03	5 41.441	- 1.00 42.23	8
MOTA	3050	OH2	$\mathbf{T}\mathbf{A}\mathbf{W}$	S 63		19.05	5 33.37	9 60.511	1.00 28.99	8.
MOTA	3051	CH2	WAT	S 64		47.92	33.25	3 61.470	1.00 34.93	8
MOTA	3052	OH2	WAT	s 65		32.50	36.00	0 41.000		8
ATOM	3053		WAT			27.24				
ATOM	3054			s 67		5.17				8
	3055									8
MOTA						41.15			1.00 27.31	8
ATOM	3056			S 69		12.869				8
MOTA	3057			s 70		17.499			1.00 24.91	8
MOTA	3058			S 71		27.152	2 12.18	9 53.999	1.00 18.76	8
MOTA	3059	OH2	WAT	s 72		25,213	54.80	9 67.866	1.00 61.35	8
ATOM	3060	OH2	WAT	s 73		17.671	48.51	5 53.188	1.00 37.63	8
ATOM	3061	OH2	WAT	S 74		23.765			1.00 21.81	8
ATOM	3062			s 75	•	35.535			1.00 34.04	
ATOM	3063		_	s 76		26.280				8
MOTA	3064			s 77		18.451			1.00 32.20	8
									1.00 28.55	8
MOTA	3065			S 78		10.446			1.00 44.74	8
ATOM	3066	OH2		5 79		13.256			1.00 35.45	8
MOTA	3067	OH2		S 80		23.571		2 69.937	1.00 49.49	8
ATOM	3068	OH2	WAT :	5 81		29.891	18.07	l 46.109	1.00 22.84	8
MOTA	3069	OH2	WAT :	82		12.886	42.723	75.807	1.00 35.31	8
MOTA	3070	OH2	WAT !	s 83 ·		41.348			1.00 47.24	8
ATOM	3071		WAT S			13.406			1.00 49.67	
ATOM	3072		WAT S			30.444	35.217		1.00 38.15	8
ATOM	3073	OH2				5.217				8
	3074						40.817		1.00 19.51	8
MOTA		CH2				8.891	21.532		1.00 30.72	8
ATOM	3075		WAT S			41.816	25.022		1.00 22.92	8
ATOM	3076		WAT S			50.621	36.644		1.00 29.29	8
MOTA	3077		WAT S			26.008	34.532		1.00 45.42	8
ATOM	3078	OH2	WAT S	91		8.131	39.168	54.903	1.00 31.50	8
ATOM	3079	OH2 V	WAT S	92		16.591	58.091	57.551	1.00 34.73	8
ATOM	3080	OH2 V	WAT S	93		34.773	54.065	69.382	1.00 36.05	8
MOTA	3081	OH2 V	WAT S	94		42.105	31.720		1.00 35.49	8
MOTA	3082	OH2 V				29.684	52.077	_	1.00 35.17	8
ATOM	3083		VAT S	_	•-	26.411	37.426		1.00 41.68	
ATOM	3084		VAT S		•	41.183	52.989			8
	3085	OH2 V							1.00 50.77	8
MOTA						21.167	6.202		1.00 33.36	8
ATOM	3086	OH2 V				25.060	18.985		1.00 46.63	8
ATOM	3087	OH2 W	VAT S	100		37.304	39.027	73.722	1.00 25.99	8
MOTA	3088	OH2 W	IAT S	101		15.911	54.635	39.343	1.00 29.88	8
ATOM	3089.	CH2 W	AT S	102 -		48.730	25.803	59.572	1.00 37.97	8
MOTA	3090	OH2 W	AT S	103		24.029	42.997	74.111	1.00 25.23	8
ATOM	3091	OH2 W				42.477	21.773	46.986	1.00 49.05	8
ATOM	3092	OH2 W				29.984	22.945	31.397	1.00 44.21	8
		OH2 W				40.850				
MOTA	3093					0.050	36.936	31.885	1.00 43.26	8
MOTA	3094	OH2 W				9.750	32.487	48.823	1.00 35.71	8
MOTA	3095	OH2 W				7.618	30.171	58.896	1.00 40.03	8
MOTA	3096	OH2 W				17.603	13.771	59.767	1.00 50.33	8
MOTA	3097	OH2 W				22.590	8.744	67.501	1.00 34.81	8
ATOM	3098	OH2 W	AT S	111		21.034	29.771	76.056	1.00 30.02	8
ATOM	3099	OH2 W				24.791	14.674	50.081	1.00 51.96	8
ATOM	3100	OH2 W				40.750	47.494	54.056	1.00 46.98	8
	3101	OH2 W				7.708	42.479	58.027	1.00 34.08	
ATOM	3102	OH2 W				32.375		77.566		8
MOTA	J 1 U Z		MI 3			JE.J/3	49.136		1.00 27.53	8
		•								

MOTA	3103	OH2 WAT 5 116	5.596	17.009	64.551	1.00 39.15	8
	3104	OH2 WAT S 117	20.194	5.0.998	70.563	1.00 19.73	8
ATOM		_					
ATOM	3105	OH2 WAT S 118	23.853	64.927	64.164	1.00 27.16	8
ATOM	3106	OH2 WAT S 119	9.277	43.601	46.279	1.00 32.31	8
ATOM	3107	OH2 WAT S 120	15.613	24.398	46.723	1.00 55.20	8
	3108	OH2 WAT S 121	33.110	16.122	54.229	1.00 35.91	8
MOTA				34.085	33.852	1.00 37.49	
ATOM	3109	OH2 WAT S 122	26.772				8
ATOM	3110	OH2 WAT S 123	28.654	37.783	75.829	1.00 47.30	8
ATOM	3111	OH2 WAT S 124	49.180	22.653	59.678	1.00 37.33	8
	3112	OH2 WAT S 125	20.561	27.788	65.975	1.00 67.86	8
MOTA					57.366	1.00 36.18	8
ATOM	3113	OH2 WAT S 126	34.251	13.344			
MOTA	3114	OH2 WAT S 127	49.215	36.854	48.117	1.00 33.63	8
MOTA	3115	OH2 WAT S 128	45.826	19.588	41.601	1.00 44.07	8
	.3116	OH2 WAT S 129	18.693	56.382	64.014	1.00 47.77	8
ATOM .			44.181	24.202	36.963	1.00 32.70	8
MOTA	3117						
MOTA	3118	OH2 WAT S 131	19.160	51.901	38.133	1.00 54.07	8
ATOM	3119	OH2 WAT S 132	16.904	36.558	48.679	1.00 42.21	. 8
ATOM	3120	OH2 WAT S 133	46.851	26.029	34.353	1.00 56.33	. 8
	3121	OH2 WAT S 134	3.925	41.533	68.647	1.00 45.99	8
ATOM							
ATOM	3122	OH2 WAT S 135	44.590	38.382		1.00 44.50	8
ATOM	3123	OH2 WAT S 136	6.384	19.317	71.166	1.00 28.17	8
ATOM	3124	OH2 WAT S 137	17.982	39.823	66.487	1.00 49.31	8
	3125	OH2 WAT S 138	8.317	22.286	61.863	1.00 43.42	8
ATOM			29.248	14.196	55.622	1.00 35.55	8
ATOM	3126	OH2 WAT S 139					
MOTA	3127	OH2 WAT S 140	30.377	33.180	80.320	1.00 43.94	8
ATOM	3128	OH2 WAT S 141	41.842	32.906	27.392	1.00 24.82	8
ATOM	3129	OH2 WAT S 142	33.971	3.859	64.002	1.00 41.93	8
	3130	OH2 WAT S 143	27.314	8.087	70.916	1.00 49.03	8
MOTA							
MOTA	3131	OH2 WAT S 144	4.310	39.006	64.550	1.00 32.70	8
ATOM	3132	OH2 WAT S 145	2.940	19.950	63.265	1.00 33.24	8
ATOM	3133	OH2 WAT S 146	24.134	47.625	60.121	1.00 44.24	8
ATOM	3134	OH2 WAT S 147	25.035	53.746	42.337	1.00 47.82	8
		OH2 WAT S 148	32.767	38.897		1.00 21.86	8
MOTA	3135						
MOTA	3136	OH2 WAT S 149	37.145	57.288	47.392	1.00 36.13	8
ATOM	3137	OH2 WAT S 150	25.171	18.011	32.273	1.00 38.04	8
ATOM	3138	OH2 WAT S 151	24.054	43.182	55.583	1.00 41.68	8
	3139	OH2 WAT S 152	27.686	64.936	52.937	1.00 60.62	8
ATOM			24.084	39.543	76.589	1.00 22.62	8
MOTA	3140	_					8
MOTA	3141	OH2 WAT S 154	42.110	10.159	68.662	1.00 46.98	
MOTA	3142	OH2 WAT S 155	9.675	22.905	75.335	1.00 26.45	8
ATOM	3143	OH2 WAT S 156	4.506	34.799	52.857	1.00 33.84	8
	3144	OH2 WAT S 157	32.583	35.051	76.446	1.00 36.27	8
ATOM			40.341	58.311	60.390	1.00 54.69	8
ATOM	3145						
ATOM	3146	OH2 WAT S 159	29.473	58.378	71.881	1.00 28.59	8
MOTA	3147	OH2 WAT S 160	11.829	60.543	56.138	1.00 37.67	8
ATOM	3148	OH2 WAT S 161	24.247	48.010	67.935	1.00 56.62	8
	3149	OH2 WAT S 162	ز 12.85	33.929	77.503	1,00 29.88	8
ATOM			9.49`	26.168	59.687	1.00 15.42	8
ATOM	3150	OH2 WAT S 163					
ATOM	3151	OH2 WAT S 164	27.424	16.480	38.895	1.00 36.86	8
ATOM	3152	OH2 WAT S 165	8.512	56.634	49.614	1.00 30.08	8
ATOM	3153	OH2 WAT S 166	30.721	13.394	57.919	1.00 39.47	8
-		OH2 WAT S 167	49.594	38.223	73.903	.1.00 29.50	8
MOTA	3154					1.00 38.12	8
ATOM	3155	OH2 WAT S 168	41.994	48.023	74.119		
ATOM	3156	OH2 WAT S 169	42.092	39.503	33.116	1.00 24.47	8
ATOM	3157	OH2 WAT S 170	34.547	12.749	38.054	1.00 38.65	8
		OH2 WAT S 171	15.377	60.862	50.791	1.00 32.82	8
ATOM	3158		31.854	42.110	62.950	1.00 42.43	8
MOTA	3159	OH2 WAT S 172					
ATOM	3160	OH2 WAT S 173	48.743	44.073	57.626	1.00 34.04	8
MOTA	3161	OH2 WAT S 174	8.723	50.038	42.232	1.00 32.87	8
	3162	OH2 WAT S 175	14.257	18.280	53.455	1.00 40.51	8
ATOM			31.917	37.509	53.943	1.00 40.43	8
ATOM	3163					1.00 47.97	
ATOM	3164	OH2 WAT S 177	23.921	47.029	70.642	1.00 41.31	8
ATOM	3165	OH2 WAT S 178	27.974	47.778	69.949	1.00 62.12	8
ATOM .		OH2 WAT S 179	7.850	25.093	51.345	1.00 50.13	8
	3167	OH2 WAT S 180	22.080	48.840	66.463	1.00 53.81	8
MOTA		OU2 WAR C 191	34.780	48.220	77.419	1.00 30.86	8
ATOM	3168	OH2 WAT S 181	54.,00	40.220			Ģ

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Figure 17-49

MOTA	3169		WAT		-		35.526	52.018		47.14	8
ATOM	3170	OH2			183	29.166	21.424	28.950		45.08	8
MOTA	3171	он2			184	51.175	51.545	62.599		33.88	8
MOTA	3172	OH2			185	18.520	46.208	42.323		50.85	8
MOTA	3173	он2			186	44.774	30.219	38.653		45.36	8
MOTA	3174		$\mathbf{T}\mathbf{A}\mathbf{W}$	S	187	30.770	9.460	69.837		32.44	8
MOTA	3175		\mathbf{v}		188	22.157	39.535	78.736		37.01	8
MOTA	3176		WAT		189	11.778	50.526	68.987		41.34	8
ATOM	3177	OH2			190	31.339	60.910	49.439		21.88	8
MOTA	3178	OH2			191	31.165		74.907		27.47	8
ATOM	3179	OH2			192	39.705	15.398	70.464		47.05	8
ATOM	3180	OH2		S	193	3.668	34.304	72.937		39.82	8
MOTA	3181	OH2			194	25.256	9.360	67.925		33.21	8
MOTA	3182	OH2			195	47.575	17.667	48.773		40.79	8
MOTA	3183	OH2				32.017	13.045	34.633		37.00	. 8
MOTA	3184		wat		197	35.476	7.006	64.436		49.59	´8
MOTA	3185				198	12.180	16.270	56.288		47.22	8
ATOM	3186		WAT		199	37.133	21.226	75.963		38.59	8
MOTA	3187	OH2	WAT		200	40.268	15.712	48.199		39.24	8
ATOM	3188	OH2	WAT	_	201	25.159	17.768	46.858		49.88	8
ATOM	3189	OH2			202	24.593	27.104	65.727		53.46	8
ATOM	3190		WAT		203	36.741	20.267	33.858		41.90	8.
ATOM	3191	OH2	WAT		204	10.013	53.930	47.546		48.06	8
MOTA	3192				205	22.305	16.731	54.471		27.07	8
MOTA	3193	OH2	TAW		206	47.454	34.778	74.101		47.44	8
MOTA	3194	OH2	WAT .	_	207	35.189	55.767	45.193		59.49	8
MOTA	3195	OH2			208	37.827	18.151	36.382		45.31	8
ATOM	3196		TAW		209	6.823	37.405	51.989		58.23	8
ATOM	3197		WAT		210	32.040	43.551	36.157		30.78	8
ATOM	3198		TAW		211	17.038	52.360	63.283		34.08	8
MOTA	3199		WAT			30.001	18.471			33.92	8
ATOM	3200		WAT			23.045	28.615	33.729		44.22	8
MOTA	3201		WAT			26.130	61.496	75.246		40.49	8
MOTA	3202		_		215	33.881	32.473	46.604		39.35	8
MOTA	3203				216	23.887	45.987	44.362		36.50	8
ATOM	3204				217	6.925	42.281	65.917		34.22	8
ATOM	3205	OH2	WAT	5	218	32.823	8.977	59.213	1.00	27.03	8
END											

PCT/US00/24700

					Figure 18-	·I		<u>.</u> .	
				Residue	# X	Y	Z	B Segment	ŤΒ
ATOM	1	CB	ALA A		46.726	14.971	138.208	1.00 56.80	
MOTA	2	C	ALA A		47.943	12.813	138.561	1.00 58.93	
	3	Ö	ALA		48.857		137.884	1.00 60.99	
ATOM					46.995		140.488	1.00 56.88	
ATOM	4	N	ALA A				139.052	1.00 59.41	
MOTA	5	CA	ALA A		46.801				
ATOM-	6	N	LYS 2		47.890		138.903	1.00 53.81	
ATOM	7	CA	LYS 2	A 3	48.937		138.492	1.00 53.62	
ATOM	8	CB	LYS Z	A 3	48.736		139.156	1.00 50.26	
MOTA	9	CG	LYS I	A 3	48.917	9.279	140.665	1.00 56.64	
MOTA	10	CD	LYS		48.950	7.891	141.285	1.00 57.18	
	11	CE	LYS		49.160	7.964	142.796	1.00 56.74	
MOTA					50.423		143.165	1.00 54.86	
MOTA	12	NZ	LYS A	_	49.063		136.986	1.00 49.95	
MOTA	13	C	LYS				136.248	1.00 44.34	
ATOM	14	0	LYS A		48.088				
ATOM	15	N	VAL 2		50.287		136.550	1.00 46.01	
ATOM	. 16	CA	VAL A	A 4	50.609		135.142	1.00 42.48	
ATOM	17	CB	VAL A	A 4	51.901		134.809	1.00 43.42	
ATOM	18	CG1	VAL A	A 4	52.179	10.713	133.307	1.00 39.20	
ATOM	19		VAL 2		51.773	12.186	135.310	1.00 39.34	
	20	C	VAL		50.787	8.510	134.806	1.00 38.41	
ATOM	21	õ	VAL		51.659	7.839	135.351	1.00 37.08	
MOTA			LYS		49.959		133.899	1.00 37.79	
MOTA	22	N			50.016		133.515	1.00 38.17	
MOTA	23	CA	LYS A				133.887	1.00 38.40	
MOTA	24	CB	LYS A	A 5	48.700		135.385	1.00 38.40	
MOTA	25	CG	LYS 3		48.411				
MOTA	26	CD	LYS 2		49.384		136.070	1.00 44.10	
MOTA	27	CE	LYS I		49.017		137.534	1.00 45.97	
MOTA	28	NZ	LYS 2	A 5	49.045		138.322	1.00 51.78	
ATOM	29	С	LYS I	A 5	50.275		132.030	1.00 38.31	
MOTA	30	0	LYS	_	49.992		131.201	1.00 38.13	
ATOM	31	N	LEU .		50.817	5.220	131.717	1.00 35.05	
	32	CA	LEU		51.082	4.818	130.346	1.00 31.46	
ATOM	33	CB	LEU		52.582	4.592		1.00 28.46	
MOTA					53.094		128.720	1.00 30.91	
MOTA	34	CG	LEU .		52.618	2.884		1.00 33.05	
ATOM	35		LEU				127.744	1.00 21.96	
ATOM	36		LEU .	_	52.630	5.312	130.164	1.00 30.50	
ATOM	37	С	LEU .	_	50.307	3.512			
MOTA	38	0	LEU 2		50.453	2.581	130.955	1.00 32.82	
MOTA	39	N	ILE A	A 7	49.459	3.456		1.00 26.94	
ATOM	40	CA	ILE 2	A 7	48.676	2.255	128.893	1.00 28.29	
ATOM	41	CB	ILE :	A 7	47.218	2.598	128.493	1.00 28.94	
ATOM	42	CG2	ILE .	a 7	46.499	1.343	128.041	1.00 32.57	
ATOM	43	CG1			46.447	3.172	129.688	1.00 36.59	
ATOM	44	CD1			46.979	4.468	130.236	1.00 46.80	
	45	C	ILE	_	49.341	1.470		1.00 31.09	
MOTA			IL :	_	49.600	2.009		1.00 27.65	
MOTA	46	0	·GL:		49.638	0.201		1.00 27.30	
ATOM	47					-0.514	127.016	1.00 25.50	
MOTA	48	CA	GL'		50.277 50.578	2 024	127.480	1.00 30.66	
MOTA	49	C	GLY .			-2.024	128.592	1.00 30.02	
MOTA	50	0	GLY A				120.334	1.00 28.94	
ATOM	51	N	THR	A 9	51.238	-2.777			
MOTA	52	CA	THE.	A 9	51.614	-4.156		1.00 33.63	
ATOM	53	CB	THR	A 9	50.393	-5.083	126.857	1.00 36.19	
ATOM	54	OG1	THR		50.827	-6.441	126.992	1.00 34.87	
	55	CG2	THR		49.633	-4.931	125.548	1.00 36.49	
MOTA	56	C	THR		52.567	-4.637	125.794	1.00 34.83	
MOTA					52.545	-4.133	124.677	1.00 36.91	
MOTA	57	0	THR		53.407	-5.609	126.129	1.00 39.15	
MOTA	58	N	LEU .					1.00 40.21	
ATOM	59	CA	LEU A		54.345	-6.167		1.00 42.40	
MOTA	60	CB	LEU 2		55.402	-7.009			
ATOM	61	CG	LEU .		56.482	-6.282		1.00 42.29	
ATOM	62	CD1	LEU .	A 10	55.870	-5.293	127.647	1.00 42.92	
ATOM	63	CD2	LEU A		57.319	-7.306	127.424	1.00 40.29	
ATOM	64	c	LEU .		53.591	-7.039	124.159	1.00 41.70	
	65	ō	LEU		54.055	-7.266	123.044	1.00 37.13	
ATOM	66	N	ASP .		52.419	-7.519	124.557	1.00 47.28	
MOTA	00	**			-	. = = -			

									_						
MOTA	67	ÇA	ASI	? A	11		51	.617	-8	.369	123.	683	1.00	53.	30
ATOM	68	CB	ASI) A	11		50	.230	-8	.608	124.	287	1,00	52.	35
	69							.295		.331					
ATOM		. CG	ASI		11								1.00		
MOTA	70	OD:	l ASI	? A	11		51	.004	-10	.358	125.0	685	1.00	52.	21
MOTA	71	OD:	2 ASI	Δ	11		49	.630	-8	.883	126.	567	1.00	5.8	48
									7	.840	120.				
MOTA	72	С	ASI		11			.459					1.00	53.	33
ATOM	73	0	ASI	A	11		51	.360	-8	.626	121.3	311	1.00	54.	31
MOTA	74	N	TYF		12			.424		.521			1.00		
MOTA	75	CA	TYF	A S	12			. 275		.970		_	1.00	51.	41
MOTA	76	CB	TYF	A 5	12		51	.328	-4	.437	120.7	755	1.00	49.	05
ATOM	77	CG	·TYF		12		50	.164	-3	.729	121.4	121	1.00		
MOTA	78	CD:	LTYF	A	12			.296		.157		986	1.00	47.	Q8
ATOM	79	CE1	TYR	A	12		49	.252	-2	.430	123.2	263	1.00	47.	53
ATOM	80	CD2		Δ.	12		48	.952	-3	. 565	120.7	749	1.00		
ATOM	81	CE2	TYR	. А	12			.906		.847			1.00		
ATOM	82	CZ	TYR	A	12		48	.061	-2	. 279	122.5	566	1.00	48.	67
MOTA	83	ОН	TYR		12		47	.030			123.1		1.00		
ATOM	84	C	TYR	. А	12			.367			119.8		1.00	50.	0.1
ATOM	85	0	TYR	. A	12		52	.197	-6.	. 525	118.5	596	1.00	45.	56
ATOM	86	N	GLY		13		53	. 484	-6	931	120.3	396	1.00	Λ Ω	72
MOTA	87	CA	GLY		13			.574		.458			1.00		
ATOM	88	C	GLY	Α	13		54	.196	-8.	.727	118.8	357	1.00	53.	32
ATOM	. 89	0	GLY	Δ	13		54	.931	-9	.184	117.9	982	1.00	52	64
MOTA	90	N	LYS		14			.045		.294	119.2		1.00		
MOTA	91	CA	LYS	Α	14		52.	. 555	-10.	.518	118.5	579	1.00	54.	56
ATOM	92	CB	LYS	A	14		52	.022	-11.	475	119.6	553	1.00	58.	02
•			LYS					086					1.00		
MOTA	93	CG			14				-12.		120.5				
ATOM	94	CD	LYS	Α	14		53.	934	-13.	154	119.9	918	1.00	61.	61
ATOM	95	CE	LYS	A	14		54.	747	-12.	638	118.7	734	1.00	61.	77
ATOM	96	NZ	LYS		14			514	-13.		118.0		1.00		
MOTA	97	С	LYS	A	14			455	-10.		117.5	59	1.00		
ATOM	98	0	LYS	Α	14		50.	911	-11.	145	116.9	42	1.00	51.	71
ATOM	99	N	TYR		15		51	143	-8	955	117.3	172	1.00	46	92
MOTA	100	CA	TYR		15			091			116.4		1.00		
ATOM	101	CB	TYR	Α	15		48.	959	-7.	915	117.2	253	1.00	50.	40
ATOM	102	CG	TYR	Α	15		48.	456	-8.	793	118.3	86	1.00	53.	01
								166		255	119.6		1.00		
MOTA	103	CD1	TYŖ		15										
MOTA	104	CE1	TYR	Α	15		47.	722	-9.	053	120.6	85	1.00	51.	72
ATOM	105	CD2	TYR	A	15		48.	283	-10.	166	118.2	808	1.00	54.	67
	106	CE2			15			838	-10.		119.2		1.00		
ATOM			TYR												
ATOM	107	CZ	TYR	A	15		47.	561	-10.	412	120.4	85	1.00	54.	18
MOTA	108	OH	TYR	Α	15		47.	130	-11.	208	121.5	20	1.00	55.	42
ATOM	109	С	TYR		15			592		617	115.3		1.00		
ATOM	110	0	TYR	Α	15			933		635	115.0			43.	
ATOM	111	N	ARG	Α	16		51.	758	-7.	924	114.7	91	1.00	46.	29
ATOM	112	CA	ARG		16		52	347		109	113.7	27	1.00	45.	66
								-					1.00		
ATOM	113	CB	ARG		16			779		545	113.4				
ATOM	114	CG	ARG	A	16		54.	677	-7.	698	114.6	36	1.00	56.	90
MOTA	115	CD	ARG	Α.	16		54.	992	-6.	388	115.3	15	1.00	60.	72
	116		ARG					021		602	116.3		1.00		
MOTA		NE			16										
MOTA	117	cz	ARG	A	16		57.	211	-7.	141	116.0	70	1.00	66.	68
MOTA	118	NH1	ARG	Α	16		57.	520	-7.	519	114.8	34 .	1.00	65.	68
	119				16		58.				117.0		1.00		
ATOM		NH2	ARG												
MOTA	120	С	ARG	Α	16		51.	573	-7.	298	112.4	29	1.00	44.	20
MOTA	121	0	ARG	Α	16		50.	871	-8.	293	112.2	54	1.00	43.4	41
	122	N				•	51.		-6.		111.5		1.00		
ATOM			TYR		17										
MOTA	123	CA	TYR	Α	17		51.		-6.	453	110.2			38.	
ATOM	124	CB	TYR	Α	17		50.	913	-5.	072	109.5	65	1.00	33.8	83
	125	CG					49.		-4.		110.0		1.00		
MOTA			TYR		17										
ATOM	126	CD1	TYR	A	17		49.		-3.		111.4		1.00		
ATOM	127	CE1	TYR		17		48.	540	-3.	184	111.9	09	1.00	27.3	16
	128	CD2	TYR		17		48.		-3.		109.2		1.00		
MOTA												_			
ATCM	129		TYR		17		47.		-2.	-	109.6		1.00		
MOTA	130	CZ	TYR	Α	17		47.	626	-2.	659	111.0	09	1.00	27.2	28
ATCM	131	ОН	TYR		17		46.		-1.		111.4		1.00	22.0)4
					_						109.3		1.00		
ATOM	132	С	TYR	A	17		51.	716	-7.	טכנ	103.5°	00	4.00	47.	- 4

	477	_				5 505 100 con	
MOTA	133		TYR A		53.150	-7.525 109.683	
MOTA	134	N	PRO A	. 18	51.440	-7.925 108.278	1.00 46.68
ATOM	135	CD	PRO A	. 18	50.076	-7.765 107.755	1.00 47.16
MOTA	136		PRO A		52.205		
MOTA	137	CB	PRO A	. 18	51.213	-9.091 106.262	1.00 48.14
ATOM	138	CG	PRO A	. 18	50.343	-7.837 106.274	1.00 55.13
ATOM	139		PRO A		53.556	-8.303 106.885	
ATOM	140		PRO A		53.788	-7.101 106.766	
MOTA	141	N	LYS A	. 19	54.432	-9.261 106.592	1.00 53.22
ATOM	142	CA	LYS A	19	55.800	-9.044 106.114	1.00 57.00
ATOM	143	CB	LYS A			-10.242 105.252	
MOTA	144		LYS A			-10.929 104.537	
ATOM	145	CD	LYS A	. 19	54.239	-9.963 103.714	1.00 70.76
MOTA	146	CE	LYS A	19	53.004	-10.653 103.162	1.00 73.70
ATOM	147	NZ	LYS A		52.116	-9.701 102.442	
MOTA	148	С	LYS A		56.229	-7.757 105.405	
ATOM	149	0	LYS A	19	` 57.230	-7.150 105.796	1.00 59.86
ATOM	150	N	asn a	20	55.515	-7.338 104.367	1.00 49.62
ATOM	151	CA	ASN A	20	55.925	-6.130 103.652	
ATOM	152	CB	ASN A	20	55.829	-6.359 102.143	
MOTA	153	CG	ASN A	20	56.729	-7.487 101.670	1.00 51.26
ATOM	154	OD1	ASN A	20	57.948	-7.437 101.843	1.00 46.88
ATOM	155	ND2		20	56.130	-8.513 101.074	
ATOM	156	С	asn a	20	55.16 7	-4.862 104.023	1.00 45.50
ATOM	157	0	ASN A	20	55.481	-3.778 103.533	1.00 45.35
ATOM	158	N	HIS A	21	54.182	-4.997 104.899	1.00 37.46
ATOM	159	CA	HIS A	21	53.374	-3.863 105.321	1.00 32.39
ATOM	160	CB	HIS A	21	52.198	-4.355 106.162	1.00 29.34
ATOM	161	CG	HIS A	21	51.118	-3.339 106.348	1.00 30.50
ATOM	162	CD2	HIS A	21	50.999	-2.314 107.223	1.00 22.88
ATOM	163		HIS A	21	49.993	-3.298 105.552	1.00 30.15
							1.00 30.96
ATOM	164		HIS A	21	49.226		
ATOM	165	NE2	HIS A	21	49.814	-1.680 106.945	1.00 36.41
ATOM	166	С	HIS A	21	54.194	-2:879 106.155	1.00 29.18
ATOM	167	0	HIS A	21	55.030	-3.279 106.963	1.00 26.92
	168	N	PRO A	22	53.965	-1.572 105.969	1.00 31.12
MOTA							
ATOM	169	CD	PRO A	22	53.027	-0.912 105.043	1.00 29.46
MOTA	170	CA	PRO A	22	54.702	-0.567 106.739	1.00 29.27
MOTA	171	CB	PRO A	22	54.012	0.732 106.326	1.00 26.00
ATOM	172	CG	PRO A	22	53.670	0.434 104.875	1.00 31.52
	173	c	PRO A	22	54.624	-0.822 108.253	1.00 29.96
ATOM							
ATOM	174	0	PRO A	22	55.575	-0.538 108.981	1.00 27.47
ATOM	175	N	LEU A	23	53.501	-1.371 108.715	1.00 26.64
MOTA	176	CA	LEU A	23	53.309	-1.644 110.144	1.00 30.44
	177	CB	LEU A	23	51.833	-1.428 110.515	1.00 24.09
MOTA							
MOTA	178	CG	LEU A	23	51.356	0.029 110.479	1.00 25.30
ATCM	179	CD1	LEU A	23 ·	49.836	0.103 110.668	1.00 17.72
ATOM	180	CD2	LEU A	23	52.086	0.816 111.574	1.00 24.15
	181	C	LEU A	23	53.775	-3.015 110.662	1.00 31.64
MOTA							
ATCM	182	0	LEU A	23	53.252		
ATOM	183	N	LYS A	24	54.753	-3.636 110.012	1.00 28.25
ATOM	184	CA	LYS A	24	55.200	-4.929 110.513	1.00 30.90
ATOM	185	CB	LYS A	24	55.718	-5.810 109.372	1.00 36.59
MOTA	186	CG	LYS A	24	57.178	-5.650 108.982	1.00 40.77
MOTA	187	CD	LYS A	24	57.546	-4.259 108.535	1.00 44.61
ATOM	188	CE	LYS A	24	· 58.858	-4.303 107.755	1.00 50.44
ATOM	189	NZ	LYS A	24	59.959	-4.990 108.487	1.00 51.30
			LYS A		56.282	-4.736 111.581	1.00 32.57
ATOM	190	С		24			
MOTA	191	0	LYS A	24	56.695	-5.683 112.245	1.00 29.83
ATOM	192	N	ILE A	25	56.729	-3.497 111.750	1.00 27.06
ATOM	193	CA	ILE A	25	-57.755	-3.200 112.739	1.00 30.45
	194	CB	ILE A	25	58.416	-1.822 112.499	1.00 33.37
ATOM						1 757 111 100	
ATOM	195	CG2		25	59.056	-1.757 111.120	1.00 33.22
ATCM ·	196	CG1	ILE A	25	57.361	-0.722 112.662	1.00 30.45
ATCM	- 197		ILE A	25	57.930	0.689 112.700	1.00 33.12
	198	C	ILE A	25	57.156	-3.129 114.141	1.00 32.10
ATCM	200	_	^	ر ب	27.120	-J.12J 447.441	2.00 30.40

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Figure 18-4

ATOM	199	0	ILE A	25		55.967	-2.851	114.310	1.00 28.15
MOTA	200		PRO A			57.979		115.168	1.00 31.64
MOTA	201					59.395		3 115.139	1.00 31.11
ATOM	202					57.507		116.556	1.00 31.04
MOTA	203					58.709		117.347	1.00 32.41
MOTA	204					59.454		116.324	1.00 39.33
	205		PRO A			57.265		116.827	1.00 28.42
MOTA	205		-			58.001		116.315	1.00 28.42
MOTA		-	PRO A			56.251		117.614	1.00 24.15
MOTA	207		ARG A						
MOTA	208					55.977		117.899	1.00 28.53
ATOM	209				-	54.787		117:048	1.00 29.77
ATOM	210	CG CD	ARG A	27 27		55.075 53.918		3 114.620	1.00 29.64 1.00 26.61
MOTA	211	NE	ARG A ARG A	27		53.510		114.520	
ATOM	212 213	CZ	ARG A	27		52.649		. 115.173	1.00 28.52 1.00 29.70
ATOM	213		ARG A 1 ARG A	27		51.857		115.173	1.00 29.70
MOTA	214		2 ARG A	27		52.451		114.983	1.00 30.17
MOTA	215		ARG A	27		55.746		119.387	1.00 23.25
MOTA		C		27				120.113	
MOTA	217	0	ARG A			56.679			1.00 24.60
MOTA	218	N	VAL A	28		54.529		119.863	1.00 23.51
MOTA	21,9	CA	VAL A	28 28		54.282 52.800		121.282	1.00 29.33
MOTA	220 221	CB	VAL A					121.033	1.00 34.56
ATOM			l VAL A	28		52.599			1.00 32.42
MOTA	222	CG2		28		51.947 55.158		120.903	1.00 33.77
ATOM	223	C	VAL A	28		55.673			1.00 29.75
MOTA	224	0	VAL A	28 29		55.341	-2.059	123.182	1.00 32.49 1.00 26.09
ATOM	225	N	SER A SER A			56.162	-2.982		1.00 26.09
ATOM	226	CA	SER A	29 29	•	56.058	-4.399		1.00 31.39
MOTA	227 228	CB OG	SER A	29		56.562		120.579	1.00 28.92
MOTA	229	C	SER A	29		57.609		122.453	1.00 34.77
MOTA	230	o	SER A	29		58.378		123.391	1.00 29.39
MOTA MOTA	231	И	LEU A	30		57.967		121.380	1.00 23.33
ATOM	232	CA	LEU A	30		59.317		121.240	1.00 32.03
ATOM	232	CB	LEU A	30		59.554		119.829	1.00 32.03
ATOM	234	CG	LEU A	30		61.008		119.333	1.00 33.22
ATOM	235		LEU A	30		61.066		118.224	1.00 28.76
ATOM	236		LEU A	30		61.948		120.441	1.00 35.11
MOTA	237	c	LEU A	30		59.423		122.236	1.00 30.29
ATOM	238	ō	LEU A	30		60.397		122.984	1.00 27.69
ATOM	239	N	LEU A	31		58.408		122.232	1.00 27.38
MOTA	240	CA	LEU A	31		58.372		123.126	1.00 24.94
ATOM	241	CB	LEU A	31		57.008		123.042	1.00 24.92
ATOM	242	CG	LEU A	31		56.918		123:460	1.00 30.49
ATOM	243	CD1		31		55.492		123.881	1.00 24.71
ATOM	244	CD2		31		57.851	4.355	124.603	1.00 27.32
ATOM	245	C	LEU A	31		58.610	1.429	124.564	1.00 28.18
ATOM	246	0	LEU A	31		59.489	1.928	125.263	1.00 33.64
ATOM	247	N	LEU A	32		57.831	0.445	125.000	1.00 30.17
ATOM	248	CA	LEU A	32		57.965	-0.084	126.357	1.00 30.59
ATOM	249	CB	LEU A	32		56.944		126.601	1.00 30.55
ATOM	250	CG	LEU A	32		55.458		126.402	1.00 29.50
MOTA	251	CD1	LEU A	32		54.611	-2.107	126.727	1.00 28.31
ATOM	. 252	CD2	LEU A	32		55.058	0.273	127.287	1.00 31.92
ATOM	253	С	LEU A	32		59.376	-0.597	126.657	1.00 33.56
MOTA	254	0	LEU A	32		59.961	-0.243	127.682	1.00 36.51
MOTA	255	N	ARG A	33		59.926			1.00 29.75
ATOM	256	CA	ARG A	33 -		61.271	-1.953	125.999	1.00 33.49
ATOM	257	CB	ARG A	33		61.630		124.945	1.00 39.50
ATOM	258	CG	ARG A	33		60.814	-4.283	125.024	1.00 44.40
MOTA	259	CD	ARG A	33		61.237	~5.256	123.933	1.00 53.68
ATOM	260	NE	ARG A	33		60.515	-6.522	124.007	1.00 56.66
ATOM	261	CZ	ARG A	33		60.611	-7.384	125.014	1.00 58.73
ATOM	262		ARG A	33		61.402	-7.121	126.045	1.00 59.32
ATOM	263		ARG A	33		59.911	-8.511	124.991	1.00 57.91
ATOM	264	С	ARG A	33	•	62.314	-0.845	125.978	1.00 31.45

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Figure 18-5

MOTA	265	O ARG	A 33		63.288	-0.88	5 126.722	1 00	26.49
ATOM	266	N PHE	A 34		62.103				
MOTA	267	CA PHE					6 125.123		32.42
					63.042	1.25		1.00	33.37
ATOM	268		A 34		62.617	2.18	0 123.858	1.00	31.68
ATOM	269		A 34		63.653	3.20	2 123.486	1.00	29.05
ATOM	270	CD1 PHE	A 34		64.825	2.81			
ATOM	271				63.458		6 123.781		
ATOM	272	CE1 PHE			65.793				28.25
ATOM	273					3.76		1.00	29.35
-			A. 34		64.416		1 123.435		
MOTA	274	CZ PHE			65.589	5.10	8 122.783	1.00	29.08
ATOM	275	C PHE	A 34		63.083	2.04	2 126.305		33.47
ATOM	276	O PHE	A 34		64.155	2.29			27.49
ATOM	277	N LYS	A 35	•	61.912	2.43			
ATOM	278	CA LYS			61.848	3.19			29.34
ATOM	279	CB LYS							31.48
					60.406	3.57		1.00	30.82
ATOM	.280	CG LYS A			59.803	4.55	2 127.395	1.00	32.98
ATOM	281	CD LYS A	35		58.404	4.97	127.790		40.93
ATOM	282	CE LYS A	35		57.410	3.82			44.56
ATOM	283	NZ LYS A	35		57.754	2.65			
ATOM	284	C LYS A			62.443			1.00	
ATOM	285					2.387		1.00	
					63.136	2.933		1.00	32.01
ATOM	286	N ASP A			62.180	1.086	129.190	1.00	36.28
ATOM	287	CA ASP A	36		62.710	0.233	130.242		37.93
ATOM	288	CB ASP A	. 36		62.145	-i.178			41.27
ATOM	289	CG ASP A			62.731	-2.117			
ATOM	290	OD1 ASP A			62.660			1.00	
ATOM	291	OD2 ASP A				-1.793		1.00	
					63.261	-3.178		1.00	45.78
ATOM	292	C ASP A			64.227	0.181	. 130.174	1.00	38.74
ATOM	293	O ASP A			64.902	0.187	131.201	1.00	36.23
MOTA	294	N ALAA	37		64.760	0.127	128.958	1.00	37.96
MOTA	295	CA ALA A	37	•	66.201	0.080			
ATOM	296	CB ALA A			66.525	-0.158			39.49
ATOM		C ALA A	37						39.74
					66.832	1.386		1.00	40.09
ATOM		O ALA A	37		67.962	1:402	129.714	1.00	38.80
ATOM		n met a	38		66.085	2.477	129.131	1.00	39.04
ATOM	300	CA MET A	38		66.567	3.789	129.545	1.00	38.71
ATOM	301	CB MET A	38		65.965	4.863		1.00	36.66
ATOM	302	CG MET A	38		66.335	4.744			
ATOM		SD MET A	38		68.005			1.00	39.16
ATOM		CE MET A				5.298			37.55
			38		67.892	7.033	127.287	1.00	35.74
ATOM		C MET A	38		66.187	4.094	130.995	1.00	40.58
ATOM		O MET A	38		66.484	5.173	131.502	1.00	38.12
ATOM	307 1	A MEA V	39		65.530	3.147	131.657	1.00	
MOTA	308 (CA ASN A	39		65.094	3.346	133.039		42.46
ATOM	309	CB ASN A	39		66.298	3.494	133.979		
ATOM		G ASN A	39						46.06
ATOM		DD1 ASN A			67.125		134.074		51.69
			39		66.625	2.175	134.487	1.00	
MOTA		JD2 ASN A	39		68.396	2 313	133.695	1.00	49.13
MOTA	313 (3 <i>9</i>		64.222	4.594	133.134	1.00	
ATOM	314 0) ASN A	39		64.375	5.402	134.050	1.00	42 74
ATOM	315 N	LEU A	40		63.301		132.188	1.00	
ATOM		A LEU A	40		62.427			1.00	40.22
							132.170	1.00	
ATOM		B LEU A	40		62.524		130.812	1.00	
ATOM		G LEU A	40		63.940	7.074	130.447	1.00	40.40
ATOM ·		D1 LEU A	40		63.916	7.753	129.088	1.00	32.75
ATOM	320 C	D2 LEU A	40		64.470		131.513	1.00	
ATOM	321 C		40		60.967		132.505		
ATOM	322 0		40		60.076			1.00	
							132.213	1.00	
ATOM			41		60.720		133.124	1.00	
MOTA	324 C		41		59.363	4.109	133.520	1.00	42.43
MOTA	325 C	B ILE A	41		58.536	3.575	132.330	1.00	
ATOM	326 C	G2 ILE A	41		59.137		131.820	1.00	
MOTA		G1 ILE A	41		57.082	3 367	132.774	1.00	
MOTA		D1 ILE A	41						
					56.147		131.676	1.00	
MOTA	329 C	ILE A	41		59.376		134.619	1.00 4	
ATCM	330 o	ILE A	41	1	60.255	2.195	134.654	1.00 4	13.05

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MOTA	331 N ASP A 42	58.414 3.148 135.532 1 00 47 8
ATOM	332 CA ASP A 42	58.301 2.183 136.620 1.00 49.2
ATOM	333 CB ASP A 42	
MOTA	334 CG ASP A 42	59.493 3.688 138.284 1.00 52.6
ATOM	335 OD1 ASP A 42	
		60.614 3.146 138.141 1.00 49.2
ATOM	336 OD2 ASP A 42	59.355 4.866 138.678 1.00 52.4
MOTA	337 C ASP A 42	
		57.034 1.368 136.405 1.00 51.2
MOTA	338 O ASP A 42	56.048 1.866 135.864 1.00 48.0
MOTA		
		57.072 0.111 136.832 1.00 51.4
MOTA	340 CA GLU A 43	55.945 -0.792 136.673 1.00 50.6°
ATOM		
		56.234 -2.094 137.412 1.00 54.49
MOTA	342 CG GLU A 43	55.208 -3.178 137.185 1.00 60.55
· ATOM		FF F0.4
		55.524 -4.432 137.974 1.00 66.12
MOTA	344 OE1 GLU A 43	54.761 -5.417 137.861 1.00 70.33
MOTA		56.536 -4.427 138.711 1.00 67.39
MOTA	346 C GLUA 43	54.645 -0.178 137.178 1.00 50.20
ATOM	347 O GLUA 43	53.567 -0.475 136.658 1.00 48.27
MOTA	348 N LYS A 44	54.755 0.683 138.186 1.00 49.04
MOTA		53.601 1.356 138.778 1.00 47.56
ATOM	350 CB LYS A 44	54.013 2.004 140.112 1.00 54.73
		= 2.00 24./3
ATOM	351 CG LYS A 44	53.190 3.229 140.542 1.00 58.07
ATOM	352 CD LYS A 44	
MOTA	353 CE LYS A 44	52.849 5.727 140.151 1.00 61.93
MOTA	354 NZ LYS A 44	
MOTA	355 C LYS A 44	52.929 2.387 137.875 1.00 44.52
ATOM	356 O. LYS A 44	F4 8F6
MOTA		53.674 2.915 136.914 1.00 41.03
MOTA	358 CA GLU A 45	
ATOM		
	359 CB GLU A 45	54.271 4.810 135.500 1.00 38.52
MOTA	360 CG GLU A 45	54.973 5.572 136.589 1.00 40.30
		54.575 5.572 136.565 1.00 40.30
MOTA		56.241 6.222 136.096 1.00 38.06
ATOM	362 OE1 GLU A 45	57.170 5.478 135.715 1.00 36.93
		57.270 5.476 135.715 1.00 36.93
MOTA	363 OE2 GLU A 45	56.306 7.467 136.084 1.00 32.57
MOTA	364 C GLUA 45.	
	- · · · · · · · · · · · · · · · · · · ·	
ATOM .	365 O GLUA 45	51.783 3.907 134.015 1.00 39.77
ATOM	366 N LEUA 46	52.700 1.953 134.645 1.00 36.90
MOTA	367 CA LEU A 46	52.165 1.207 133.517 1.00 40.46
ATOM	368 CB LEU A 46	53.222 0.219 133.034 1.00 35.52
ATOM	369 CG LEU A 46	52.873 -0.619 131.811 1.00 43.75
ATOM	370 CD1 LEU A 46	52.571 0.292 130.630 1.00 42.06
ATOM	371 CD2 LEU A 46	54.035 -1.544 131.500 1.00 42.90
MOTA	372 C LEUA 46	
ATOM	373 O LEUA 46	50.741 -0.306 134.730 1.00 39.37
ATOM	374 N ILE A 47	
MOTA	375 CA ILE A 47	48.560 0.068 133.033 1.00 32.12
ATOM	376 CB ILE A 47	
ATOM	377 CG2 ILE A 47	46.069 0.360 132.833 1.00 30.60
ATOM	378 CG1 ILE A 47	47.448 2.015 134.156 1.00 36.56
ATOM	379 CD1 ILE A 47	46.372 3.080 134.162 1.00 35.46
ATOM	380 C ILE A 47	48.428 -0.920 131.882 1.00 33.67
ATOM	381 O ILE A 47	48.505 -0.532 130.717 1.00 27.64
ATOM	382 N LYS A 48	48.231 -2.195 132.205 1.00 32.98
ATOM	383 CA LYS A 48	48.102 -3.224 131.176 1.00 30.98
ATOM	384 CB LYS A 48	
MOTA	385 CG LYS A 48	47.956 -5.747 130.819 1.00 46.81
ATOM	386 CD LYS A 48	
ATOM	387 CE LYS A 48	47.967 -8.240 130.492 1.00 54.43
	388 NZ LYS A 48	
ATOM		
ATOM	389 C LYS A 48	46.869 -3.006 130.310 1.00 29.55
ATCM		45.764 -2.840 130.820 1.00 29.65
ATOM	391 N SERA 49	47.071 -2.992 128.996 1.00 30.69
ATOM	392 CA SER A 49	45.989 -2.802 128.033 1.00 29.32
MOTA	393 CB SER A 49	46.551 -2.805 126.609 1.00 31.53
ATOM	394 OG SER A 49	47.571 -1.834 126.443 1.00 30.74
ATOM	395 C SER A 49	44.952 -3.916 128.147 1.00 31.31
atom	396 O SERA 49 '	45.295 -5.059 128.436 1.00 34.44
	•	

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Figure 18-7

					2 502 105 000	1 00 00 00
ATOM	397			43.688	-3.582 127.922	1.00 32.87
ATOM	398	CA ABG	A 50	42.632	-4.582 127.960	1.00 31.45
MOTA	399	CB ARG	A 50	41.636	-4.325 129.101	1.00 28.35
ATOM	400	CG ARG	A 50	40.729	-3.103 128.915	1.00 32.05
	401			39.653	-3.055 130.008	1.00 30.46
MOTA						1.00 25.21
MOTA	402			38.821	-1.850 129.964	
MOTA	403	CZ ARG		37.930	-1.569 129.016	1.00 28.32
ATOM	404	NH1 ARG	A 50	37.726	-2.406 128.001	1.00 25.45
ATOM	405	NH2 ARG	A 50	37.238	-0.439 129.087	1.00 24.92
MOTA	406	•		41.894	-4.470 126.638	1.00 31.12
	407		_	41.895	-3.406 126.019	1.00 24.62
MOTA				41.264	-5.566 126.181	1.00 32.55
MOTA	408	N PRO				
MOTA	409	CD PRO I		41.164	-6.921 126.751	1.00 32.40
ATOM	410	CA PRO A	4 51	40.534	-5.506 124.917	1.00 30.36
ATOM	411	CB PRO A	A 51	40.138	-6.967 124.683	1.00 33.95
ATOM	412	CG PRO A	A 51	41.173	-7.750 125.499	1.00 32.85
	413	C PRO		39.309	-4.630 125.134	1.00 31.61
ATOM				38.877	-4.431 126.267	1.00 29.84
MOTA	414	O PRO				
MOTA	415	N ALA A		38.755	-4.093 124.058	1.00 29.09
ATOM	416	CA ALA		37.556	-3.294 124.183	1.00 29.61
ATOM	417	CB ALA A	A 52	37 <i>.</i> 365	-2.447 122.956	1.00 28.67
ATOM	418	C ALA A	52	36.437	-4.321 124.288	1.00 32.39
	419	O ALA A		36.603	-5.453 123.844	1.00 30.40
ATOM	420	N THR		35.318	-3.947 124.896	1.00 32.98
ATOM						1.00 36.61
ATOM	421	CA THR A		34.192		
ATOM	422	CB THR A		33.253	-4.514 126.166	1.00 34.22
MOTA	423	OG1 THR A		32.734	-3.193 125.970	1.00 29.52
ATOM	424	CG2 THR A	53	33.998	-4.579 127.493	1.00 36.45
ATOM	425	C THR A	53	33.411	-4.700 123.702	1.00 38.94
	426	O THR A		33.559	-3.689 123.012	1.00 32.67
ATOM	427	N LYS A		32.577	-5.679 123.372	1.00 39.19
MOTA				31.792	-5.595 122.152	1.00 40.71
MOTA	428					
MOTA	429	CB LYS A		30.933	-6.851,121.994	1.00 41.68
ATOM	430	CG LYS A		30.367	-7.034 120.597	1.00 49.42
ATOM	431	CD LYS A	54	29.541	-8.310 120.508	1.00 51.82
ATOM	432	CE LYS A	54	29.075	-8.588 119.087	1.00 52.94
MOTA	433	NZ LYS A	54	30.216	-8.879 118.182	1.00 54.26
	434	C LYS A		30.913	-4.347 122.237	1.00 39.46
ATOM	435	O LYS A		30.719	-3.637 121.249	1.00 37.19
MOTA					-4.075 123.434	1.00 36.71
ATOM	436	N GLU A		30.404		
MOTA	437	CA GLU A		29.554	-2.913 123.665	1.00 36.18
MOTA	438	CB GLU A		29.109	-2.877 125.127	1.00 42.16
ATOM	439	CG GLU A	55	28.223	-1.694 125.476	1.00 46.04
ATOM	440	CD GLU A	. 55	27.873	-1.639 126.953	1.00 51.15
ATOM	441	OE1 GLU A		27.092	-0.748 127.343	1.00 56.53
	442	OE2 GLU A		28.382	-2.482 127.727	1.00 51.67
ATOM	443	C GLU A		30.278	-1.607 123.32:	1.00 35.45
MOTA					-0.729 122.66	1.00 29.11
MOTA	444	O GLU A		29.721		1.00 27.82
ATOM	445	Й GLU У		31.518	-1.476 123.776	
ATOM	446	CA GLU A		32.289	-0.269 123.497	1.00 34.70
ATOM	447	CB GLU A	56	33.635	-0.329 124.232	1.00 30.71
ATOM	448	CG GLU A	56	33.474	-0.484 125.746	1.00 35.09
	449	CD GLU A		34.787	-0.675 126.479	1.00 32.29
MOTA	450	OE1 GLU A		35.645	-1.434 125.986	1.00 34.54
ATOM					-0.094 127.569	1.00 33.25
MOTA	451	OE2 GLU A		34.951		
ATOM	452	C GLU A		32.495	-0.104 121.988	1.00 32.51
ATOM	453	O GLU A		32.341	0.990 121.444	1.00 29.59
ATOM	454	N LEU A	57	32.827	-1.196 121.311	1.00 35.58
	455	CA LEU A	57	33.039	-1.147 119.871	1.00 35.70
ATOM	456	CB LEU A		33.475	-2.512 119.334	1.00 35.25
MOTA				34.829	-3.030 119.814	1.00 36.19
ATOM	457				-4.390 119.183	1.00 33.69
ATOM	458	CD1 LEU A		35.095		
MOTA	459	CD2 LEU A		35.925	-2.041 119.433	1.00 32.61
ATOM	460	C LEU A	57	31.772	-0.717 119.157	1.00 36.02
ATOM	461	O LEU A	57	31.828	0.067 118.205	1.00 32.72
	462	N LEU A	_	30.631	-1.228 119.620	1.00 32.35
ATCM			-			

MOTA	463 CA LEU A 58	20 252	0 000 110 001	
		29.353	-0.898 119.004	1.00 33.21
MOTA		28.260	-1.844 119.495	1.00 35.17
ATOM	465 CG LEU A 58	28.504	-3.296 119.077	1.00 33.71
ATOM	466 CD1 LEU A 58	27.338	-4.166 119.524	1.00 36.80
MOTA	467 CD2 LEU A 58	28.665	-3.364 117.570	1.00 36.50
MOTA	468 C LEUA 58	28.940	0.543 119.222	1.00 30.99
MOTA	469 O LEU A 58	27.915	0.985 118.700	1.00 35.50
ATOM	470 N LEU A 59	29.733	1.279 119.993	
ATOM	471 CA LEU A 59	29.443		1.00 32.55
MOTA	472 CB LEU A 59		2.687 120.217	1.00 30.37
ATOM		30.387	3.279 121.268	1.00 28.01
		30.174	2.828 122.716	1.00 32.19
ATOM	474 CD1 LEU A 59	31.248	3.427 123.604	1.00 24.85
MOTA	475 CD2 LEU A 59	28.785	3.263 123.192	1.00 25.65
ATOM	476 C LEU A 59	29.632	3.405 118.890	1.00 31.26
MOTA	477 O LEUA 59	29.020	4.442 118.652	1.00 31.80
MOTA	478 N PHE A 60	30.482	2.850 118.026	1.00 29.79
ATOM	479 CA PHE A 60	30.726	3.454 116.716	1.00 30.24
ATOM	480 CB PHE A 60	32.131	4.055 116.637	1.00 29.99
ATOM	481 CG PHE A 60	32.443	4.691 115.299	1.00 28.88
ATOM	482 CD1 PHE A 60	31.706	5.780 114.845	
ATOM	483 CD2 PHE A 60			1.00 25.58
ATOM		33.448	4.178 114.479	1.00 24.00
		31.959	6.351 113.592	1.00 26.12
ATOM	485 CE2 PHE A 60	33.709	4.740 113.226	1.00 25.98
ATOM	486 CZ PHE A 60	32.963	5.828 112.781	1.00 24.53
MOTA	487 C PHE A 60	30.536	2.520 115.529	1.00 30.30
ATOM	488 O PHE A 60	29.810	2.854 114.602	1.00 32.82
ATOM	489 N HIS A 61	31.195	1.363 115,543	1.00 32.85
MOTA	490 CA HIS A 61	31.075	0.418 114.431	1.00 34.59
ATOM	491 CB HIS A 61	32.296	-0.492 114.361	1.00 32.89
ATOM	492 CG HIS A 61	33.576	0.238 114.116	1.00 34.25
ATOM	493 CD2 HIS A 61	34.225	0.532 112.967	1.00 34.23
ATOM	494 ND1 HIS A 61	34.328	0.786 115.133	
MOTA	495 CE1 HIS A 61	35.390		
ATOM	496 NE2 HIS A 61	35.350		1.00 37.50
ATOM			1:243 113.307	1.00 37.91
		29.824	-0.449 114.480	1.00 38.44
ATOM	498 O HIS A 61	29.213	-0.612 115.538	1.00 35.78
MOTA	499 N THR A 62	29.462	-1.015 113.327	1.00 39.73
ATOM	500 CA THR A 62	28.278	-1.868 113.218	1.00 38.05
MOTA	501 CB THR A 62	27.682	-1.825 111.804	1.00 37.22
ATOM	502 OG1 THR A 62	28.631	-2.345 110.867	1.00 41.15
MOTA	503 CG2 THR A 62	27.348	-0.404 111.418	1.00 38.27
ATOM	504 C THR A 62	28.598	-3.317 113.551	1.00 39.06
ATOM ·	505 O THR A 62	29.731	-3.768 113.392	1.00 39.32
ATOM	506 N GLUA 63	27.582	-4.034 114.017	1.00 40.92
ATOM	507 CA GLU A 63	27.696	-5.441 114.393	1.00 40.68
MOTA	508 CB GLU A 63	26.303	-6.000 114.704	1.00 43.19
MOTA	509 CG GLU A 63	26.269	-7.451 115.171	1.00 46.90
ATOM	510 CD GLU A 63	26.472		
MOTA	511 QE1 GLU A 63			
ATOM				1.00 52.78
			-6.556 117.358	1.00 57.24
ATOM	513 C GLU A 63		-6.263 113.268	1.00 36.19
MOTA	514 O GLUA 63		-7.011 113.481	1.00 29.70
ATOM	515 N ASP A 64	27.755	-6.119 112.074	1.00 35.85
ATOM	516 CA ASP A 64	28.198	-6.841 110.886	1.00 37.61
ATOM	517 CB ASP A 64	27.363	-6.382 109.697	1.00 43.30
MOTA	518 CG ASP A 64	27.313	-4.872 109.582	1.00 53.38
MOTA	519 OD1 ASP A 64	28.290	-4.269 109.089	1.00 52.15
ATOM	520 OD2 ASP A 64	26.298	-4.285 110.018	1.00 53.97
ATOM	521 C ASP A 64		-6.660 110.594	1.00 35.04
	522 0 ASP A 64		-7.625 110.303	
MOTA				1.00 33.60
ATOM	523 N TYR A 65		-5.423 110.671	1.00 33.88
ATOM	524 CA TYR A 65		-5.153 110.419	1.00 32.91
ATCM	525 CB TYR A 65		-3.637 110.375	1.00 34.80
ATOM	526 CG TYR A 65		-3.253 110.219	1.00 35.19
ATOM	527 CD1 TYR A 65	34.009 -	-3.751 109.163	1.00 28.43
ATOM	528 CE1 TYR A 65	35.352 -	-3.411 109.024	1.00 32.52

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                                    33.863
                                            -2.398 111.134
                                                              1.00 34.08
         530
               CE2 TYR A
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                                           -2.050 111.002
                                                              1.00 29.89
MOTA
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ATOM
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                   TYR A
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                                                                   35.29
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                                    37.286
MOTA
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               OH
                   TYR A
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                           65
                                    32.405
                                                              1.00 27.65
MOTA
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                   TYR A
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                                    33.339
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                                                              1.00
MOTA
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                                                   111.209
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                                                              1.00 27.32
ATOM
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                   ILE
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                                            -6.153 113.858
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ATOM
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               CA
                   ILE A
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         537
                           66
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                                            -5.764 115.217
                                                              1.00 32.25
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                                    33.029
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MOTA
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MOTA
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         552
              CA
                   THR A
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MOTA
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                   THR A
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MOTA
ATOM
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              С
                   THR A
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         557
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                   THR A
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                          70
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ATOM
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ATOM
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MOTA
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              CE
                  MET A
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                                                              1.00 37.36
ATOM
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                  MET A
                                   36.484 -12.685 113.401
ATOM
                                   37.392 -13.488
                                                   113.607
                                                              1.00 31.47
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                  MET A
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ATOM
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                                                              1.00 36.6
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                                                   111.681
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                                   34.245 -14.160
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                  GLU A
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MOTA
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         577
              CG
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                                   34.206 -15.359
                                                   109.897
MOTA
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                                   34.257 -16.693 110.633
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ATOM
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                          71
         580
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                  GLU A
MOTA
                  GLU A
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                                   36.732 -14.169 110.769
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MOTA
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                                                                   32.99
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                  GLU A
                          71
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ATOM
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ATOM
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                  ALA A
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                  ALA A
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                                   38.225 -12.981 109.264
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              CA
MOTA
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                                   38.366 -11.580
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              CB
                  ALA A
ATOM
                          72
                                   39.498 -13.362 109.998
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         586
              С
                  ALA A
ATOM
                                                             1.00
                                                                   31.53
                                   40.337 -14.094 109.466
                  ALA A
                          72
ATOM
         587
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                          73
                                   39.647 -12.873
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                                                                   30.87
                  GLU A
                                                   111.224
         588
              N
ATOM
                                   40.847 -13.177
                                                   111.985
                                                             1.00 29.66
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              CA
                  GLU A
MOTA
                                                             1.00 33.33
                          73
                                   41.004 -12.224 113.180
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              CB
                  GLU A
ATOM
                                   42.234 -12.545
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                  GLU A
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              CG
ATOM
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                                   42.390 -11.634 115.233
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        592
              CD
                  GLU A
ATOM
                                                             1.00 41.02
                                   42.601 -10.418 115.044
                          73
        593
              OE1
                  GLU A
MOTA
        594
              OE2 GLU A
                                   42.298 -12.138 116.372
                                                             1.00 41.21
MOTA
```

MOTA	595	С	GLU A	73	40.906 -14.615 112.485 1.00 31.73
				73	41.957 -15.249 112.409 1.00 32.96
ATOM	596	0	GLU A		
MOTA	597	N	ARG A	74	39.798 -15.145 112.992 1.00 35.85
ATOM	598	CA	ARG A	74	39.847 -16.511 113.502 1.00 43.24
-	599	CB	ARG A	74	38.548 -16.892 114.216 1.00 43.63
MOTA					
ATOM	600	CG	ARG A	74	37.450 -17.349 113.294 1.00 51.20
ATOM	601	CD	ARG A	74	36.366 -18.087 114.063 1.00 51.13
	602	NE	ARG A	74	35.534 -18.871 113.158 1.00 57.40
ATOM					
MOTA	603	CZ	ARG A	. 74	35.991 -19.870 112.403 1.00 56.36
ATOM	604	NH1	ARG A	74	37.273 -20.208 112.446 1.00 51.10
	605	NH2		74	35.172 -20.517 111.586 1.00 58.75
MOTA					
ATOM	606	С	ARG A	74	40.125 -17.506 112.372 1.00 43.06
ATOM	607	0	ARG A	74	40.916 -18.429 112.541 1.00 42.52
	608	N	SER A	75	39.485 -17.305 111.222 1.00 43.63
MOTA					
ATOM	609	CA	SER A	75	
MOTA	610	CB	SER A	75	38.485 -18.089 109.113 1.00 42.05
ATOM	611	OG	SER A	75	38.420 -16.799 108.532 1.00 38.43
					40.910 -17.797 109.282 1.00 46.44
ATOM	612	С	SER A	75	
ATOM	613	0	SER A	75	41.339 -18.522 108.383 1.00 45.17
ATOM	614	N	GLN A	76	41.466 -16.638 109.618 1.00 46.18
					42.642 -16.116 108.936 1.00 44.73
ATOM	615	CA	GLN A	76	
ATOM	616	ÇВ	GLN A	76	43.868 -16.973 109.226 1.00 37.36
ATOM	617	CG	GLN A	76	45.162 -16.208 109.045 1.00 43.96
				76	45.415 -15.214 110.176 1.00 41.86
MOTA	618	CD	GLN A		
ATOM	619	OE1	GLN A	76	44.499 -14.537 110.655 1.00 37.78
ATOM	620	NE2	GLN A	76	46.669 -15.111 110.591 1.00 45.00
MOTA	621	С	GLN A	76	42.374 -16.120 107.429 1.00 44.17
MOTA	622	0	GLN A	76	
ATOM	623	N	SER A	. 77	41.168 -15.713 107.053 1.00 43.13
ATOM	624	CA	SER A	77	40.784 -15.667 105.649 1.00 44.66
					40.182 -17.004 105.220 1.00 44.56
MOTA	625	CB	SER A	77	
ATOM	626	OG	SER A	77	38.974 -17.246 105.925 1.00 42.58
ATOM	627	С	SER A	77	39.747 -14.573 105.448 1.00 44.80
				77	39.096 -14.142 106.395 1.00 45.11
ATOM	628	0	SER A		
ATOM	629	N	VAL A	78	39.590 -14.137 104.207 1.00 46.06
ATOM	630	CA	VAL A	78	38.632 -13.095 103.888 1.00 47.65
	631	CB	VAL A	78	39.107 -12.245 102.701 1.00 49.63
ATOM					
ATOM	632	CG1		78	
ATOM	633	CG2	VAL A	78	40.454 -11.627 103.017 1.00 53.00
ATOM'	634	С	VAL A	78	37.275 -13.682 103.530 1.00 48.07
				78	37.111 -14.301 102.480 1.00 42.31
MOTA	635	0	VAL A		
MOTA	636	N	PRO A	79	36.282 -13.492 104.407 1.00 49.82
MOTA	637	CD	PRO A	79	36.347 -12.782 105.696 1.00 50.83
	638	CA	PRO A	79	34.927 -13.998 104.186 1.00 51.33
MOTA					34.170 -13.450 105.396 1.00 53.13
ATOM	639	CB	PRO A	79	· · · · · · · · · · · · · · · · · · ·
MOTA	640	CG	PRO A	-9	35.244 -13.469 106.469 1.00 53.50
ATOM	641	С	PRO A	; 9	34.343 -13.517 102.858 1.00 52.42
				. 9	34.670 -12.428 102.382 1.00 55.73
MOTA	642	0	PRO A		
ATOM	643	N	LYS A	80	33.482 -14.343 102.273 1.00 49.63
ATOM	644	CA	LYS A	80	32.824 -14.053 101.002 1.00 51.62
	645	CB	LYS A	80	31.632 -15.004 100.822 1.00 53.92
MOTA					30.817 -14.808 99.545 1.00 56.27
MOTA	646	CG	LYS A	80	
MOTA	647	CD	LYS A	80	29.586 -15.712 99.560 1.00 56.61
	648	CE	LYS A	80	28.744 -15.579 98.298 1.00 56.04
MOTA					29.471 -16.036 97.081 1.00 58.90
MOTA	649	NZ	LYS A	80	
ATOM	650	С	LYS A	80	32.338 -12.607 100.874 1.00 51.10
MOTA	651	0	LYS A	80	31.539 -12.140 101.689 1.00 49.22
			GLY A	81	32.821 -11.914 99.842 1.00 51.14
MOTA	652	N			
MOTA	653	CA	GLY A	81	
ATOM	654	C	GLY A	81	32.876 ÷9.496 100.599 1.00 46.90
	655	ō	GLY A	81	32.671 -8.301 100.397 1.00 43.90
ATOM					33.504 -9.942 101.681 1.00 44.50
ATOM	656	Ŋ	ALA A	82	
MOTA	657	CA	ALA A	82	33.973 -9.029 102.715 1.00 44.69
	658	CB	ALA A	82	34.497 -9.825 103.903 1.00 44.62
ATOM				82	35.049 -8.073 102.215 1.00 41.82
MOTA	659	C	ALA A		
ATOM	660	0	ALA A	82	
					•

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Figure 18-11

ATOM	661	N ARG	A 83	35.874	-8.549 101.289	1.00 43.30
ATOM	662	CA ARG	A 83	36.959	-7.742 100.741	1.00 43.25
ATOM	663	CB ARG	A 83	37.715	-8.533 99.677	1.00 46.60
ATOM	664	CG ARG		38.988		
						1.00 51.32
MOTA	665	CD ARG		39.636	-8.632 98.086	1.00 55.55
MOTA	666	NE ARG	A 83	40.995	-8.164 97.810	1.00 64.08
ATOM	667	CZ ARG	A 83	41.330	-6.905 97.540	1.00 63.01
ATOM	668					
		NH1 ARG		40.403	-5.954 97.504	1.00 62.76
MOTA	669	NH2 ARG		42.599	-6.600 97.304	1.00 59.66
ATOM	670	C ARG 2	83 A	36.453	-6.435 100.134	1.00 44.58
MOTA	671	O ARG		37.002	-5.365 100.395	
						1.00 38.05
ATOM	672	N GLU		35.404	-6.528 99.323	1.00 41.82
ATOM:	673	CA GLU A	A 84	34.824	~5.356 98.678	1.00 41.44
MOTA	674	CB GLU A	A 84	34.145	-5.765 97.367	1.00 46.27
MOTA	675	CG GLU A		33.621	-7.185 97.388	
MOTA	676					1.00 52.61
		CD GLU A		34.749	-8.198 97.308	1.00 54.12
MOTA	677	OE1 GLU A	4 84	34.555	-9.344 97.764	1.00 59.66
ATOM	678	OE2 GLU A	A 84	35.823	~7.850 96.769	1.00 50.30
ATOM	679	C GLU A	84	33.831	-4.595 99.545	
	680					1.00 37.36
ATOM		O GLU A		33.692	-3.379 99.416	1.00 34.30
MOTA	681	N LYS A		33.138	-5.301 100.427	1.00 36.00
ATOM	682	CA LYS A	85	32.154	-4.646 101.280	1.00 36.95
ATOM	683	CB LYS A		31.089	-5.649 101.725	1.00 36.60
ATOM	684	CG LYS A				
				29.975	-5.042 102.570	1.00 40.72
ATOM	685	CD LYS A		28.939	-6.092 102.963	1.00 46.21
ATOM	686	CE LYS A	85	27.839	-5.487 103.827	1.00 49.06
MOTA	687	NZ LYS A	85	26.859	-6.513 104.287	1.00 52.72
MOTA	688	C LYS A		32.785		
						1.00 36.48
atom	689	O LYS A		32.353	-2.949 102.966	1.00 32.97
MOTA	690	N TYR A	86	33.819	-4.649 103.041	1.00 33.69
ATOM	691	CA TYR A	86	34.468	-4.169 104.250	1.00 35.23
ATOM	692	CB TYR A		34.410	-5.281 105.300	
					75.281 105.300	1.00 33.65
MOTA	693	CG TYR A		32.990	-5.665 105.680	1.00 35.09
MOTA	694	CD1 TYR A		. 32.165	-4.765 106.351	1.00 34.06
ATOM	695	CE1 TYR A	86	30.866	-5.100 106.704	1.00 34.32
MOTA	696	CD2 TYR A		32.470	-6.923 105.365	1.00 33.17
ATOM	697	CE2 TYR A				
				31.162	-7.271 105.716	1.00 33.91
MOTA	698	CZ TYR A		30.369	-6.350 106.386	1.00 34.21
MOTA	699	OH TYR A	86	29.079	-6.658 106.738	1.00 35.20
MOTA	700	C TYR A	86	35.901	-3.672 104.046	1.00 36.09
ATOM	701	O TYR A		36.552	-3.208 104.984	
						1.00 36.06
ATOM	702	N ASN A	87	36.382	-3.777 102.814	1.00 36.46
MOTA	703	CA ASN A	87	37.712	-3.313 102.441	1.00 32.71
ATOM	704	CB ASN A	87	37.768	-1.791 102.576	1.00 36.26
MOTA	705	CG ASN A	87	38.989	-1.199 101.926	1.00 37.25
ATOM	706	OD1 ASN A				
			87	39.305	-1.518 100.784	1.00 36.29
MOTA	707	ND2 ASN A	87	39.675	-0.320 102.640	1.00 45.25
ATOM	708	C ASN A	87	38.855 ⁻	-3.956 103.217	1.00 34.73
ATOM	709	O ASN A	87	39.868	-3.315 103.512	1.00 33.23
ATOM	710	N ILE A	88	38.687	-5.237 103.523	
						1.00 32.48
ATOM	711	CA ILE A	88	39.676	-6.018 104.248	1.00 33.65
ATOM	712	CB ILE A	88	39.030	-6.732 105.445	1.00 38.66
ATOM	713	CG2 ILE A	88	40.021	~7.680 106.081	1.00 41.31
	714	CG1 ILE A	88	38.536	-5.707 106.461	
ATOM						1.00 40.45
ATOM	715	CD1 ILE A	88	39.641	-4.953 107.124	1.00 42.25
ATOM	716	C ILE A	88	40.251	~7.090 103.318	1.00 37.36
ATOM	717	O ILE A	88	39.555	-7.587 102.431	1.00 35.47
		N GLY A	89	41.517	-7.446 103.520	
ATOM						1.00 31.52
ATOM		CA GLY A	89	42.124	-8.477 102.698	1.00 33.53
MOTA	720	C GLY A	89	43.134	-7.994 101.675	1.00 34.50
ATOM		O GLY A	89	43.951	-8.777 101.186	1.00 32.81
		N GLY A	90	43.071	-6.710 101.335	
ATOM						1.00 31.39
MOTA		CA GLY A	90	44.005	-6.158 100.371	1.00 23.90
ATOM		C GLY A	90	45.340	-5.893 101.040	1.00 28.78
MOTA	725	O GLY A	90	45.563	-6.339 102.163	1.00 21.71
		N TYR A	91	46.221	-5.155 100.367	
ATOM	.20	14 TYV V	71	40.00	-J.1JJ 100.36/	1.00 28.26

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MOTA	727 CA TYR A 91	47.539 -4.850 100.918	1.00 27.34
MOTA	728 CB TYR A 91		
MOTA		48.066 -3.039 99.194	1.00 24.28
ATOM	730 CD1 TYR A 91	48.374 -1.829 99.822	1.00 21.55
MOTA	731 CE1 TYR A 91	47.970 -0.609 99.275	
MOTA	732 CD2 TYR A 91		1.00 24.05
MOTA		46.931 -1.786 97.447	1.00 29.92
MOTA	734 CZ TYR A 91	47.250 -0.597 98.086	
MOTA	735 OH TYR A 91	46.861 0.593 97.516	
ATOM	736 C TYR A 91		
MOTA	737 O TYR A 91	. 48.314 -3.689 102.869	1.00 27.20
MOTA	738 N GLUA 92	46.402 -2.971 101.938	
MOTA	739 CA, GLU A 92	46.232 -1.879 102.882	1.00 28.38
ATOM	740 CB GLU A 92		
			1.00 28.57
MOTA	741 CG GLU A 92	45.232 0.471 102.982	1.00 36.94
MOTA	742 CD GLU A 92	44.178 1.395 102.396	1.00 37.40
ATOM	743 OE1 GLU A 92	42.999 1.293 102.794	1.00 31.22
ATOM	744 OE2 GLU A 92	44.527 2.209 101.516	
			1.00 40.54
MOTA		45.770 -2.343 104.259	1.00 29.20
MOTA	746 O GLUA 92	46.389 -2.015 105.268	1.00 21.86
ATOM	747 N. ASNA 93	44.687 -3.117 104.286	1.00 26.51
MOTA	748 CA ASN A 93	44.109 -3.613 105.527	
ATOM			1.00 24.02
		42.727 -2.988 105.690	1.00 24.51
MOTA	750 CG ASN A 93	42.738 -1.488 105.405	1.00 28.61
ATOM	751 OD1 ASN A 93	43.428 -0.727 106.079	1.00 25.30
MOTA	752 ND2 ASN A 93	41.987 -1.063 104.393	1.00 20.45
ATOM	753 C ASN A 93		
			1.00 24.79
ATOM	754 O ASN A 93	42.905 -5.680 105.291	1.00 21.89
ATOM	755 N PRO A 94	45.142 -5.828 105.429	1.00 24.60
MOTA	756 CD PRO A 94	46.493 -5.246 105.540	1.00 22.93
ATOM	757 CA PRO A 94		
			1.00 27.23
ATOM		46.730 -7.488 105.093	1.00 25.46
MOTA	759 CG PRO A 94	47.299 -6.431 106.046	1.00 26.20
ATOM	760 C PRO A 94	44.743 -8.112 106.489	1.00 31.04
ATOM	761 O PRO A 94	44.411 -7.589 107.558	1.00 29.10
ATOM	762 N VAL A 95		
	_		1.00 28.27
MOTA	_	44.299 -10.367 107.291	1.00 28.82
MOTA	764 CB VAL A 95	43.938 -11.737 106.677	1.00 30.75
MOTA	765 CG1 VAL A 95	43.745 -12.766 107.776	1.00 33.60
ATOM	766 CG2 VAL A 95	42.679 -11.611 105.849	
ATOM	767 C VAL A 95		1.00 24.87
		45.503 -10.549 108.204	1.00 29.98
ATOM	768 0 VAL A 95	46.637 -10.649 107.729	1.00 31.36
ATOM	769 N SER A 96	45.264 -10.572 109.510	1.00 29.38
ATOM	770 CA SER A 96	46.335 -10.766 110.485	1.00 32.56
ATOM	771 CB SER A 96	47.325 -9.600 110.454	
			1.00 34.15
MOTA		46 758 -8.448 111.051	1.00 28.33
MOTA	773 C SER A. 96	45.681 -10.804 111.854	1.00 32.10
MOTA	774 G SERA 96	44 458 -10.839 111.950	1.00 37.91
ATOM	775 N TYRA 97	46.484 -10.795 112.913	1.00 32.57
ATOM	776 CA TYR A 97		
			1.00 34.95
MOTA		46.685 -11.735 115.182	1.00 35.47
ATOM	778 CG TYR A 97	46.492 -13.187 114.817	1.00 40.65
ATOM	779 CD1 TYR A 97	47.319 -13.812 113.882	1.00 40.63
MOTA	780 CE1 TYR A 97	47.083 -15.121 113.475	
		45 403 13 12 113 475	1.00 42.16
ATOM		45.421 -13.910 115.338	1.00 38.82
ATOM	782 CE2 TYR A 97	45.175 -15.219 114.936	1.00 42.82
ATOM	783 CE TYR A 97	46.010 -15.816 114.005	1.00 42.56
ATOM	784 OH TYR A 97	45.772 -17.105 113.601	
	785 C TYR A 97		1.00 46.03
MOTA			1.00 37.56
atom	786 O TYR A 97	45.601 -9.195 115.998	1.00 39.06
ATOM '	787 N ALA A 98	46.115 -8.418 113.948	1.00 31.96
ATOM	788 CA ALA A 98		1.00 30.43
ATOM	789 CB ALA A 98	48 405	
			1.00 29.64
ATOM			1.00 30.35
ATOM	791 O ALA A 98	44.099 -5.655 114.612	1.00 31.82
ATOM	792 N MET A 99		1.00 30.40

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Figure 18-13

	50B	
ATOM	793 CA MET A .99	42.788 -6.730 112.420 1.00 27.54
ATOM	794 CB MET A 99	42.370 -7.620 111.238 1.00 30.55
ATOM	795 CG MET A 99	42.052 -9.071 111.565 1.00 31.59
ATOM	796 SD MET A 99	
ATOM	797 CE MET A 99	
ATOM-		40.770 -9.086 109.085 1.00 28.02
		41.703 -6.696 113.490 1.00 28.02
ATOM	799 O MET A 99	40.818 -5.842 113.446 1.00 24.53
MOTA	800 N PHE A 100	41.752 -7.614 114.449 1.00 26.07
ATOM	801 CA PHE A 100	40.759 -7.583 115.516 1.00 30.47
ATOM	802 CB PHE A 100	39.738 -8.718 115.404 1.00 30.29
ATOM	803 CG PHE A 100	
ATOM	804 CD1 PHE A. 100	
	005 CD1 PME A. 100	1.00 27.01
ATOM	805 CD2 PHE A 100	38.756 -9.506 117.575- 1.00 30.68
MOTA	806 CE1 PHE A 100	36.834 -7.507 117.519 1.00 31.41
MOTA	807 CE2 PHE A 100	37.873 -9.356 118.644 1.00 28.39
MOTA	808 CZ PHE A 100	36.913 -8.355 118.618 1.00 24.06
MOTA	809 C PHE A 100	41.345 -7.616 116.922 1.00 29.67
ATOM	810 O PHE A 100	41.028 -6.751 117.740 1.00 29.67
ATOM	811 N THR A 101	
ATOM	812 CA THR A 101	
		42.770 -8.701 118.562 1.00 31.37
MOTA	813 CB THR A 101	43.610 -9.977 118.732 1.00 31.63
ATOM	814 OG1 THR A 101	42.777 -11.119 118.532 1.00 31.64
MOTA	815 CG2 THR A 101	44.197 -10.045 120.137 1.00 27.38
MOTA	816 C THR A 101	43.647 -7.493 118.884 1.00 31.66
ATOM	817 O THR A 101	43.502 -6.875 119.942 1.00 30.71
ATOM	818 N GLY A 102	44.562 -7.166 117.976 1.00 27.40
ATOM	819 CA GLY A 102	
ATOM	820 C GLY A 102	
		44.631 -4.728 118.266 1.00 27.26
ATOM	821 O GLY A 102	44.785 -3.940 119.201 1.00 27.68
MOTA	822 N SER A 103	43.767 -4.515 117.279 1.00 30.52
ATOM	823 CA SER A 103	42.941 -3.314 117.216 1.00 31.91
MOTA	824 CB SER A 103	42.085 -3.334 115.949 1.00 34.63
ATOM	825 OG SER A 103	42.896 -3.265 114.791 1.00 35.94
ATOM	826 C SER A 103	42.046 -3.163 118.441 1.00 32.44
ATOM	827 O SER A 103	
ATOM	828 N SER A 104	41.891 -2.065 118.984 1.00 25.78
		41.455 -4.270 118.871 1.00 30.47
ATOM		40.584 -4.251 120.038 1.00 30.22
ATOM	830 CB SER A 104	39.978 -5.633 120.265 1.00 23.88
ATOM	831 OG SER A 104	39.078 -5.595 121.358 1.00 36.91
ATOM	832 C SER A 104	41.367 -3.841 121.282 - 1.00 28.13
ATOM	833 O SER A 104	40.872 -3.098 122.130 1.00 25.16
ATOM	834 N LEU A 105	42.594 -4.336 121.386 1.00 29.39
ATOM	835 CA LEU A 105	43.445 -4.034 122.530 1.00 29.52
ATOM	836 CB LEU A 105	44.684 -4.922 122.471 1.00 32.90
ATOM	837 CG LEU A 105	
ATOM	838 CD1 LEU A 105	
		44.520 -5.723 124.828 1.00 35.95
MOTA	839 CD2 LEU A 105	46.582 -6.178 123.462 1.00 40.23
MOTA	840 C LEU A 105	43.834 -2.552 122.511 1.00 32.09
ATOM	841 O LEU A 105	43.896 -1.894 123.554 1.00 30.38
MOTA	842 N ALA A 106	44.081 -2.029 121.314 1.00 30.26
ATOM	843 CA ALA A 106	44.448 -0.626 121.151 1.00 28.31
ATOM	844 CB ALA A 106	44.958 -0.386 119.738 1.00 23.88
ATOM	845 C ALA A 106	43.243 0.268 121.434 1.00 26.04
ATOM	846 O ALA A 106	
ATOM	847 N THR A 107	42.058 -0.224 121.099 1.00 26.86
MOTA	848 CA THR A 107	40.841 0.542 121.322 1.00 25.04
MOTA	849 CB THR A 107	39.706 0.007 120.443 1.00 26.50
ATOM	850 OG1 THR A 107	40.111 0.092 119.069 1.00 24.62
MOTA	851 CG2 THR A 107	38.439 0.824 120.629 1.00 19.80
ATOM	852 C THR A 107	40.450 0.503 122.798 1.00 27.90
ATOM	853 O THR A 107	40.039 1.515 123.361 1.00 29.04
ATOM	854 N GLY A 108	
	855 CA GLY A 108	
ATOM		
MOTA	856 C GLY A 108	41.181 0.155 125.603 1.00 23.86
MOTA	857 O GLY A 108	40.771 0.790 126.572 1.00 26.97
ATOM	858 N SER A 109	42.434 0.236 125.158 1.00 23.07

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MOTA	859	9 CA SER A 109	43.421	1.090 125.807	1.00 20.96
ATOM	860	O CB SER A 109	44.795		1.00 24.84
ATOM	86:				
	•		45.294		1.00 25.84
ATOM	862	2 C SER A 109	43.008	2.552 125.759	1.00 21.13
MOTA	863	3 O SER A 109	43.323	3.312 126.672	1.00 23.17
ATOM	864		42.311	2.949 124.698	
					1.00 20.83
MOTA	869		41.841	4.327 124.583	1.00 21.84
ATOM	866	CB THR A 110	41.332	4.648 123.161	1.00 24.33
MOTA	867	OG1 THR A 110	42.452	4.769 122.276	1.00 25.38
ATOM	868		40.543	5.954 123.144	
					1.00 21.18
MOTA	869		40.725	4.561 125.600	1.00 28.52
MOTA	870	O THR A 110	40.632	5.637 126.197	1.00 28.27
ATOM	871	. N VAL A 111	39.882	3.558 125.809	1.00 26.88
ATOM .	872		38.811	3.706 126.793	
					1.00 30.04
MOTA	873		37.820	2.519 126.742	1.00 29.94
MOTA	874	CG1 VAL A 111	36.737	2.693 127.802	1.00 27.07
ATOM	875	CG2 VAL A 111	37.193	2.431 125.355	1.00 25.26
ATOM	876		39.440	3.797 128.187	1.00 28.10
	877				
MOTA			38.968	4.539 129.039	1.00 26.06
MOTA	878		40.521	3.056 128.415	1.00 23.92
MOTA	879	CA GLN A 112	41.188	3.097 129.711	1.00 30.27
MOTA	880	CB GLN A 112	42.268	2.020 129.804	1.00 28.61
ATOM	881				
			41.777	0.629 129.481	1.00 28.90
ATOM	882	CD GLN A 112	42.883	-0.397 129.564	1.00 28.60
MOTA	883	OE1 GLN A 112	43.344	-0.740 130.653	1.00 29.68
MOTA	884	NE2 GLN A 112	43.333	-0.880 128.409	1.00 22.13
ATOM	885	C GLN A 112	41.834	4.461 129.931	
					1.00 29.99
ATOM	886	O GLN A 112	41.791	5.006 131.035	1.00 28.43
MOTA	887	N. ALA A 113	42.453	5.004 128.885	1.00 28.64
ATOM	888	CA ALA A 113	43.083	6.315 129.001	1.00 26.62
MOTA	889	CB ALA A 113	43.693	6.732 127.684	1.00 23.49
	890				
MOTA				7.507 207.407	1.00 24.63
MOTA	891	O ALA A 113	42.232	8.183 130.240	1.00 26.38
MOTA	892	N ILE A 114	40.824	7.163 128.822	1.00 25.26
MOTA	893	CA ILE A 114	39.728	8.063 129.145	1.00 27.05
ATOM	894	CB ILE A 114	38.554	7.887 128.156	
					1.00 26.93
ATOM	895	CG2 ILE A 114	37.387	8.770 128.576	1.00 25.86
ATOM	896	CG1 ILE A 114	39.008	8.259 126.739	1.00 28.38
ATOM	897	CD1 ILE A 114	37.938	8.105 125.669	1.00 28.64
ATOM .	898	C ILE A 114	39.239	7.823 130.578	1.00 31.36
ATOM	899	O ILE A 114	38.898		
				8.770 131.291	1.00 24.56
ATOM	900	N GLU A 115	39.210	6.563 131.005	1.00 31.17
ATOM	901	CA GLU A 115	38.750	6.257 132.358	1.00 32.12
ATOM	902	CB GLU A 115	38.729	4.744 132.607	1.00 32.15
ATOM	903	CG GLU A 115	37.904	3.947 131.598	1.00 32.84
	904				
ATOM			37.875	2.459 131.912	1.00 34.12
ATOM	905	OE1 GLU A 115	38.910	1.910 132.345	1.00 30.36
MOTA	906	OE2 GLU A 115	36.826	1.827 131.699	1.00 31.38
ATOM	907	C GLU A 115	39.675	6.932 133.357	1.00 31.65
ATOM	908	O GLU A 115	39.224	7.446 134.383	1.00 29.25
ATOM	909	N GLU A 116	40.970	6.933 133.053	1.00 31.50
ATOM	910	CA GLU A 116	41.942	7.564 133.934	1.00 32.34
ATOM	911	CB GLU A 116	43.367	7.285 133.457	1.00 33.29
ATOM	912	CG GLU A 116	43.805	5.842 133.633	1.00 32.29
	913				_
MOTA			43.701	5.378 135.079	1.00 36.87
ATOM	914	OE1 GLU A 116	44.329	6.003 135.961	1.00 34.07
ATOM	915	OE2 GLU A 116	42.993	4.385 135.335	1.00 35.00
ATOM	916	C GLU A 116	41.702	9.067 134.006	1.00 36.69
ATOM	917	O GLU A 116	41.863	9.678 135.066	
					1.00 34.39
ATOM	918	N PHE A 117	41.317	9.661 132.881	1.00 31.19
ATOM	919	CA PHE A 117	41.038	11.091 132.841	1.00 28.43
ATOM	920	CB PHE A 117	40.593	11.509 131.444	1.00 30.14
	921	CG PHE A 117	40.044	12.908 131.381	
ATOM					1.00 35.78
ATOM	922	CD1 PHE A 117	40.882	14.006 131.501	1.00 33.39
ATOM	923	CD2 PHE A 117	38.675	13.123 131.225	1.00 38.90
MOTA	924	CE1 PHE A 117	40.372	15.299 131.466	1.00 30.65

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Figure 18-15

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ATOM	925	CE	2 PHE A 1	17	38.153	14.412	131.190	1.00 36.50
MOTA	926				39.003	15.501		1.00 35.41
ATOM	927	' C	PHE A 1	17	39.908	11.401	133.811	1.00 32.78
	928	0	PHE A 1		39.966	12.377		
MOTA								1.00 29.82
ATOM	929	N	LEU A 1	18	38.874	10.568	133.771	1.00 28.61
ATOM	930	CA	LEU A 1	ι Ω	37.720	10.751	134.632	1.00 32.00
MOTA	931	CB	LEU A 1	18	36.621	9.748	134.263	1.00 29.19
MOTA	932	CG	LEU A 1	8	36.098	9.830	132.820	1.00 34.47
MOTA	933	CD	l LEU A·1	18	34.962	8.836	132.622	1.00 32.69
ATOM	934	CD	2 LEU A 1	18	35.612	11.240	132.522	1.00 32.24
ATOM .	935		LEU A 1		38.123	10.590		1.00 31.17
MOTA	936	0	LEU A 1	L8	37:576	11.260	136.964	1.00 28.32
	937	N	LYS A 1	0	39.083	9.707		1.00 27.23
MOTA								
MOTA	938	CA	LYS A 1	.9	39.531	9.497	137.733	1.00 30.95
MOTA	939	CB	LYS A 1	Q	40.203	8.130	137.884	1.00 26.35
MOTA	940	CG	LYS A 1		39.293	6.954		1.00 32.44
ATOM	941	CD	LYS A 1	.9	39.895	5.624	137.986	1.00 33.31
	942	CE	LYS A 1		41.280	5.385		
MOTA								1.00 33.47
MOTA	943	NZ	LYS A 1	.9	41.874	4.102	137.904	1.00 33.40
ATOM	944	С	LYS A 1		40.493	10.594	138.173	1.00 32.65
MOTA	945	0	LYS A 1	.9	41.050	10.548	139.270	1.00 28.83
ATOM	946	N	GLY A 12	0	40.689	11.583	137.308	1.00 33.77
	947	CA	GLY A 12			12.677		
MOTA					41.571	•		1.00 33.84
ATOM	948	С	GLY A 12	0	43.035	12.448	137.340	1.00 34.27
ATOM	949	0	GLY A 12	n	43.880	13.227	137.776	1.00 36.80
MOTA	950	N	ASN A 12	1	43.347	11.384	136.606	1.00 30.77
ATOM	951	CA	ASN A 12	1	44.731	11.122	136.244	1.00 31.73
MOTA	952	CB	ASN A 12		45.089	9.646		1.00 29.34
ATOM	953	CG	ASN A 12	1	44.856	9.170	137.851	1.00 35.83
ATOM	954	OD1	ASN A 12	1	45.190	9.861	138.816	1.00 32.74
ATOM	955	ND2	ASN A 12	1	44.304	7.970	137.986	1.00 33.20
MOTA	956	С	ASN A 12	1	44.954	11.506	134.790	1.00 32.59
		O	ASN A 12	_				
MOTA	957				44.031	11.952		1.00 34.69
ATOM	958	N	VAL A 12	2	46.186	11:334	134.322	1.00 32.74
ATOM	959	CA	VAL A 12	2	46.540	11.653	132.946	1.00 33.59
ATOM	960	CB	VAL A 12	2	47.571	12.790	132.882	1.00 36.05
MOTA	961	CG1	VAL A 12	2	47.884	13.121	131.438	1.00 37.58
	962	CG2						
ATOM					47.029	14.021	133.602	1.00 37.19
ATOM	963	C	VAL A 12	2	47.147	10.397	132.352	1.00 34.47
ATOM	964	0	VAL A 12	2	48.053	9.801	132.939	1.00 31.28
MOTA	965	N	ALA A 12		46.646	9.989	131.196	1.00 28.06
ATOM	966	CA	ALA A 12	3	47.142	8.784	130.563	1.00 30.73
	967	CB	ALA A 12		46.133	7.666	130.727	1.00 32.69
MOTA								
ATOM	968	С	ALA A 12	3	47.466	8.969	129.088	1.00 30.55
ATOM	969	0	ALA A 12	3	46.909	9.830	128.406	1.00 32.89
	970	N						
MOTA			PHE A 12		48.380	8.136	128.613	1.00 27.53
MOTA	971	CA	PHE A 12	4	48.807	8.157	127.229	1.00 26.56
	972	CB	PHE A 12		50.261	8.660	127.157	1.00 25.32
ATOM								
MOTA	973	CG	PHE A 12	4	50.903	8.544	125.793	1.00 27.84
ATOM	974	CD1	PHE A 12	4	50.179	8.785	124.629	1.00 24.77
	975							1.00 21.79
ATOM		CD2	PHE A 12	4	52.266	8.266	125.686	
ATOM	976	CE1	PHE A 12	1	50.802	8.753	123.385	1.00 29.19
	977	CE2	PHE A 12		52.894		124.449	1.00 27.38
ATOM						8.235		
ATOM	978	CZ	PHE A 12	1	52.164	8.478	123.296	1.00 20.91
ATOM	979	С	PHE A 12	1	48.671	6.749	126.675	1.00 21.13
-								
MOTA	980	0	PHE A 12		49.181	5.795	127.260	1.00 25.38
ATOM	981	N	ASN A 12	5	47.933	6.624	125.580	1.00 18.87
	982	CA	ASN A 12		47.750	5.342	124.905	1.00 25.05
atom								
ATOM	983	CB	ASN A 12)	46.271	4.982	124.756	1.00,22.99
	984	CG	ASN A 12	;	46.073	3.784	123.856	1.00 24.08
ATOM								
ATOM	985		ASN A 12		46.916		123.822	1.00 20.46
ATOM	986	ND2	ASN A 12	5	44.960	3.748	123.138	1.00 16.10
	987	C	ASN A 12		48.380		123.518	1.00 23.43
MOTA								
ATOM	988	0	ASN A 12		47.718		122.542	1.00 23.48
ATOM	989	N	PRO A 12	5	49.680	5.103	123.423	1.00 24.55
						4 770	124.519	1.00 22.87
MOTA	990	CD	PRO A 12	ı	50.589	4./50	TEE. 317	1.00 22.07

MOTA	99:	L CA	PRO A 126	50.413	·5.130 122.160	1.00 22.39
ATOM	992			51.829	4.751 122.594	1.00 18.20
ATOM	993			51.564		
					3.849 123.798	1.00 25.43
MOTA	994		PRO A 126	49.867	4.224 121.058	1.00 23.18
ATOM	995	5 0	PRO A 126	50.173	4.436 119.893	1.00 20.12
ATOM	996	5 N	ALA A 127	49.058	3.232 121.423	1.00 23.27
MOTA	997	CA	ALA A 127	48.493	2.306 120.444	1.00 23.89
ATOM	998			48.176	0.967 121.118	1.00 24.82
ATOM	999		ALA A 127	47.241		
					2.864 119.778	1.00 24.76
MOTA	1000		ALA A 127	46.806	2.360 118.745	1.00 28.99
MOTA	1001		GLY A 128	- 46.666	3.906 120.367	1.00 22.12
MOTA	1002	CA	GLY A 128	45.461	4.494 119.809	1.00 21.43
MOTA	1003	С	GLY A 128	45.732	5.521 118.725	1.00 23.55
ATOM	1004	0	GLY A 128	46.875	5.695 118.291	1.00 23.25
ATOM	1005		GLY A 129	44.680	6.199 118.283	1.00 18.03
ATOM	1006		GLY A 129	44.822	7.205 117.243	_
						1.00 24.99
ATOM	1007		GLY A 129	44.600	6.655 115.847	1.00 25.11
MOTA	1008		GLY A 129	44.963	7.293 114.857	1.00 24.99
MOTA	1009	N	MET A 130	44.002	5.470 115.765	1.00 20.01
MOTA	1010	CA	MET A 130	43.729	4.825 114.481	1.00 23.63
ATOM	1011	CB	MET A 130	43.360	3.361 114.744	1.00 22.77
MOTA	1012	CG	MET A 130	44.455	2.661 115.563	1.00 26.30
ATOM	1013	SD	MET A 130	44.198	0.913 115.989	1.00 26.57
MOTA	1014	CE	MET A 130	42.665	1.030 116.936	1.00 27.59
MOTA	1015	C	MET A 130	42.580	5.617 113.869	1.00 23.70
MOTA	1016	0	MET A 130	41.421	5.199 113.901	1.00 26.28
ATOM	1017	N	HIS A 131	42.926	6.766 113.294	1.00 20.66
ATOM	1018	CA	HIS A 131	41.933	7.687 112.775	1.00 20.99
MOTA	1019	CB	HIS A 131	42.474	9.125 112.891	1.00 21.01
ATOM	1020	CG	HIS A 131	43.699	9.391 112.069	1.00 28.30
ATOM	1021		HIS A 131	44.498	8.549 111.373	1.00 19.65
	1022		HIS A 131			
MOTA				44.246	10.649 111.917	1.00 27.76
MOTA	1023		HIS A 131	45.328	10.567 111.163	1.00 20.48
MOTA	1024	NE2			9.302 110.820	1.00 24.18
ATOM	1025	С	HIS A 131	41.280	7.513 111.416	1.00 23.76
ATOM	1026	0	HIS A 131	40.453	8.341 111.051	1.00 21.95
MOTA	1027	N	HIS A 132	41.600	6.449 110.682	1.00 25.12
ATOM	1028	CA	HIS A 132	41.006	6.257 109.354	1.00 23.32
ATOM	1029	CB	HIS A 132	42.060	5.715 108.388	1.00 17.87
ATOM	1030	CG	HIS A 132	43.148	6.689 108.072	1.00 24.79
	1031		HIS A 132			
MOTA				44.496	6.574 108.144	1.00 21.72
MOTA	1032		HIS A 132	42.896	7.944 107.556	1.00 13.58
ATOM	1033		HIS A 132	44.044	8.558 107.323	1.00 15.41
MOTA	1034	NE2		45.028	7.748 107.668	1.00 15.27
ATOM	1035	С	HIS A 132	39.752	5.386 109.208	1.00 23.38
ATOM	1036	0	HIS A 132	38.947	5.615 108.304	1 00 24.70
ATOM	1037	N	ALA A 133	. 39.587	4.388 110.070	1.00 23.34
ATOM	1038	CA	ALA A 133	38.453	3.471 109.953	1 00 23.77
ATOM	1039	СВ	ALA A 133	38.515	2.417 111.053	1.00 27.49
			ALA A 133		4.145 109.966	
ATOM	1040	C		37.093		1.00 23.02
MOTA	1041	0	ALA A 133	36.878	5.117 110.691	1.00 25.98
MOTA	1042	N	PHE A 134	36.179	3.633 109.148	1.00 18.90
MOTA	1043	CA	PHE A 134	34.831	4.173 109.103	1.00 23.73
ATOM	1044	CB	PHE A 134	34.317	4.296 107.663	1.00 24.29
MOTA	1045	CG	PHE A 134	35.119	5.225 106.801	1.00 26.67
ATOM	1046		PHE A 134	36.025	4.724 105.867	1.00 28.69
	1047		PHE A 134			
MOTA				34.975	6.605 106.921	1.00 32.49
MOTA	1048		PHE A 134	36.775	5.582 105.063	1.00 28.65
ATOM	1049		PHE A 134	35.724	7.479 106.119	1.00 27.86
ATOM	1050	CZ	PHE A 134	36.623	6.967 105.188	1.00 23.93
MOTA	1051	С	PHE A 134	33.894	3.260 109.884	1.00 25.91
ATOM	1052	0	PHE A 134	34.270	2,172 110.319	1.00 27.20
MOTA	1053	N	LYS A 135	32.670	3.728 110.062	1.00 29.14
	1054	CA	LYS A 135	31.638	2.984 110.765	1.00 35.26
ATOM			LYS A 135	30.294		1.00 35.26
ATOM	1055	CB			3.628 110.429	
ATOM	1056	CG	LYS A 135	29.072	2.779 110.667	1.00 46.26

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Figure 18-17

	1057	-	7 110	A 135		27 274	2 5/2	110.211	1.00 47.72
MOTA	1057	CD	712	A 133		27.834			
MOTA	1058	CE		A 135		26.610		110.169	1.00 53.65
ATOM	1059	NZ	LYS	A 135		26.788	1.549	109.167	1.00 53.27
ATOM	1060	С		A 135		31.617	1.490	110.414	1.00 35.62
	1061			A 135		31.609		111.301	1.00 32.58
ATOM		0							
ATOM-	1062	N		A 136		31.629		109.122	1.00 35.97
MOTA	1063	CA	SER	A 136		31.555	-0.211	108.684	1.00 38.99
ATOM	1064	CB	SER	A 136		30.172	-0.474	108.083	1.00 38.87
MOTA	1065	OG		A 136		29.146	-0.072	108.975	1.00 43.54
								107.660	1.00 37.84
atom	1066	С		A 136		32.608			
ATOM	1067	0	SER	A 136		32.350		106.828	1.00 36.33
MOTA	1068	N	ARG	A 137		33.788	-0.008	107.705	1.00 33.23
MOTA	1069	CA	ARG	A 137	•	34.797	-0.368	106.724	1.00 30.89
	1070	CB		A 137		34.456		105.385	1.00 33.88
MOTA									
MOTA	1071	CG		A 137		35.009		104.201	1.00 44.41
ATOM	1072	CD	ARG	A 137		34.809		102.880	1.00 46.27
MOTA	1073	NE	ARG	A 137		35.091	-0.645	101.768	1.00 48.87
ATOM	1074	CZ		A 137		35.352	-0.261	100.526	1.00 48.64
		NH1		A 137		35.372		100.220	1.00 51.82
ATOM	1075								
ATOM	1076	NH2		A 137		35.592		99.589	1.00 49.01
ATOM	1077	С	ARG	A 137		36.209	0.021	107.143	1.00 31.84
ATOM	1078	0	ARG	A 137		36.428	1.079	107.742	1.00 30.36
ATOM	1079	Ň		A 138		37.166		106.828	1.00 30.06
								107.158	1.00 32.24
MOTA	1080	CA		A 138		38.560			
ATOM	1081	CB	ALA	A 138		39.367		107.048	1.00 31.25
ATOM	1082	С	ALA	A 138		39.095	0.449	106.187	1.00 29.49
ATOM	1083	0	ALA	A 138		38.612	0.551	105.063	1.00 26.11
	1084	N		A 139		40.099		106.615	1.00 29.54
ATOM								105.767	1.00 26.99
MOTA	1085	CA		A 139		40.673			
ATOM	1086	CB		A 139		39.685		105.662	1.00 24.10
MOTA	1087	CG	ASN	A 139		40.209	4.556	104.811	1.00 28.02
ATOM	1088	OD1	ASN	A 139		40.729	4.334	103.727	1.00 26.90
ATOM	1089			A 139		40.050		105.293	1.00 23.55
						42.027		106.285	1.00 30.17
MOTA	1090	C		A 139					
ATOM	1091	0		A 139		42.245		107.497	1.00 27.55
MOTA	1092	N	GLY	A 140		42.944		105.354	1.00 31.82
ATOM	1093	CA	GLY	A 140		44.277	3.428	105.702	1.00 24.90
ATOM	1094	Ċ		A 140		45.000	2.696	106.816	1,00 27.79
	1095	ō		A 140		45.560		107.705	1.00 23.85
ATOM								106.768	1.00 24.35
MOTA	1096	N		A 141		45.006			
ATOM	1097	CA	PHE	A 141		45.679		107.783	1.00 24.53
MOTA	1098	CB	PHE	A 141		47.031	1.146	108.197	1.00 26.40
ATOM	1099	CG		A 141		47.997	1.366	107.062	1.00 30.31
	1100			A 141		49.145		107.269	1.00 31.60
ATOM						47.781		105.802	1.00 29.44
ATOM	1101	CD2		A 141					
ATOM	1102	CE1		A 141		50.066		106.243	1.00 30.44
MOTA	1103 -	CE2	PHE	A 141		48.694	1.008	104.770	1.00 27.91
ATOM	1104	CZ	PHE	A 141		49.840	1.771	104.991	1.00 29.38
MOTA	1105	С		A 141		44.846	0.387	109.056	1.00 23.53
	1106	ō		A 141		45.194	-0.399	109.941	1.00 23.09
MOTA									1.00 22.86
ATOM	1107	N		A 142		43.760	1.143	109.159	1.00 22.00
ATOM	1108	CA	CYS .	A 142		42.925		110.356	1.00 23.87
MOTA	1109	CB	CYS .	A 142		42.472		110.723	1.00 22.51
ATOM	1110	SG		A 142		43.828	3.683	111.072	1.00 27.62
	1111	C		A 142		41.694		110.233	1.00 24.20
ATOM						40.932			1.00 24.12
MOTA	1112	Ο.		A 142			0.307		
MOTA	1113	N	TYR .	A 143		41.498	-0.663	111.219	1.00 23.84
MOTA	1114	CA		A 143		40.335		111.236	1.00 26.07
ATOM	1115	CB		A 143		40.728		111.680	1.00 27.89
				A 143		41.829	-3.582	110.855	1.00 27.30
ATOM	1116	CG							
ATOM	1117	CD1		A 143		43.169		111.137	1.00 25.76
ATOM	1118	CE1	TYR A	A 143		44.185	-3.875	110.346	1.00 25.77
MOTA	1119	CD2	TYR A	A 143		41.526	-4.394	109.762	1.00 25.87
ATOM .	1120	CE2	ጥሃፑ	A 143		42.531	-4.941	108.967	1.00 23.10
	1121	CZ		A 143		43.854	-4 679	109.262	1.00 22.93
MOTA						44.849	_5 217	108.476	1.00 20.64
ATOM	1122	OH	TYK	A 143		33.047	~5.41/	700.310	1.00 20.09

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MOTA	1123	C TYR A 143	39.281	-0.991 112.193	1.00 24.56
MCTA	1124		38.085	-1.030 111.905	
ATOM	1125	N ILE A 144	39.734	-0.471 113.331	1.00 23.77
MOTA	1126		38.833	0.102 114.335	1.00 27.11
ATOM	1127		38.871	-0.729 115.643	1.00 24.56
MOTA	1128	CG2 ILE A 144	37.941	-0.120 116.690	1.00 23.47
ATOM	1129		38.430	-2.169 115.346	1.00 28.51
MOTA	1130		38.535	-3.113 116.539	1.00 28.70
MOTA	1131	C ILE A 144	39.248	1.550 114.627	1.00 24.15
ATOM	1132		40.428	1.843 114.800	1.00 24.42
ATOM	1133		38.277	2.453 114.669	1.00 22.04
ATOM	1134	CA ASN A 145	38.555	3.866 114.920	1.00 21.31
MOTA	1135	CB ASN A 145	37.559	4.732 114.133	1.00 18.87
			37.956		
MOTA	1136	CG ASN A 145		6.205 114.091	1.00 22.21
ATOM	1137	OD1 ASN A 145	38.223	6.823 115.124	1.00 22.47
ATOM	1138	ND2 ASN A 145	37.978	6.776 112.892	1.00 23.78
ATOM	1139	C ASN A 145	38.417	4.141 116.418	1.00 22.63
MOTA	1140	O ASN A 145	37.338	4.535 116.880	1.00 22.45
ATOM	1141	N ASNA 146	39.495	3.941 117.178	1.00 16.63
MOTA	1142	CA ASN A 146	39.423	4.160 118.628	1.00 23.57
			40.708		
MOTA	1143			3.678 119.320	1.00 19.80
MOTA	1144	CG ASN A 146	41.924	4.508 118.967	1.00 27.81
MOTA	1145	OD1 ASN A 146	42.299	5.421 119.704	1.00 19.55
	1146	ND2 ASN A 146	42.544	4.202 117.827	1.00 19.55
ATOM					
MOTA	1147	C ASN A 146	39.079	5.602 119.023	1.00 26.32
MOTA	1148	O ASN A 146	38.452	5.827 120.059	1.00 28.34
ATOM	1149	N PRO A 147	39.512	6.605 118.231	1.00 28.46
			40.383	6.637 117.042	
MOTA	1150				1.00 27.18
MOTA	1151	CA PRO A 147	39.150	7.972 118.618	1.00 24.15
ATOM	1152	CB PRO A 147	39.859	8.815 117.558	1.00 25.13
MOTA	1153	CG PRO A 147	41.081	7.959 117.235	1.00 30.05
	1154	C PRO A 147	37.618	8.136 118.578	1.00 26.71
MOTA					
MOTA	1155	O PRO A 147	37.017	8.760 119.456	1.00 24.93
MOTA	1156	N ALA A 148	36.989	7:557 117.562	1.00 21.42
MOTA	1157	CA ALA A 148	35.536	7.633 117.416	1.00 21.03
	1158	CB ALA A 148	35.112	7.044 116.072	1.00 19.98
MOTA					
MOTA	1159	C ALA A 148	34.838	6.891 118.552	1.00 20.49
MOTA	1160	O ALA A 148	33.822	7.344 119.067	1.00 21.44
ATOM	1161	N VAL A 149	35.381	5.739 118.928	1.00 19.20
	1162	CA VAL A 149	34.818	4.950 120.016	1.00 24.61
MOTA	-				
ATOM	1163	CB VAL A 149	35.570	3.608 120.181	1.00 25.96
MOTA	1164	CG1 VAL A 149	35.158	2.918 121.485	1.00 26.58
ATOM	1165	CG2 VAL A 149	35.262	2.704 118.995	1.00 25.67
ATOM	1166	C VAL A 149	34.947	5.752 121.304	1.00 23.56
MOTA	1167	O VAL A 149	33.990	5.887 122.064	1.00 22.52
ATOM	1168	N GLY A 150	36.143	6.287 121.536	1.00 24.65
ATOM	1169	CA GLY A 150	36.390	7.074 122.731	1.00 22.82
	1170	C GLY A 150	35.477	8.281 122.838	1.00 25.46
ATOM					
ATOM	1171	O GLY A 150	34.919	8.564 123.904	1.00 23.17
ATOM	1172	N ILE A 151	35.327	9.001 121.733	1.00 24.38
ATOM	1173	CA ILE A 151	34.481	10.180 121.716	1.00 22.85
			34.610	10.928 120.371	1.00 24.45
ATOM	1174			10.926 120.371	
ATOM	1175	CG2 ILE A 151	33.598	12.077 120.306	1.00 24.71
ATOM	1176	CG1 ILE A 151	36.041	11.462 120.222	1.00 28.02
	1177	CD1 ILE A 151	36.354	12.056 118.854	1.00 27.10
ATOM	1170				
ATOM	1178	C ILE A 151	33.018	9.806 121.987	1.00 28.19
ATOM	1179	O ILE A 151	32.337	10.482 122.763	1.00 26.37
ATOM	1180	N GLU A 152	32.532	8.734 121.364	1.00 26.32
	1181	CA GLU A 152	31.149	8.314 121.601	1.00 30.07
ATOM					
ATOM	1182	CB GLU A 152	30.758	7.161 120.672	1.00 29.37
ATOM	1183	CG GLU A 152	30.609	7.543 119.194	1.00 27.68
ATOM	1184	CD GLU A 152	29.455	8.504 118.946	1,00 31.82
	1185	OE1 GLU A 152	29.139	8.777 117.773	1.00 33.51
ATOM					
ATOM '	1186	OE2 GLU A 152	28.862	9.009 119.918	1.00 34.73
ATOM	1187	C GLU A 152	31.009	7.879 123.055	1.00 28.00
ATOM	1188	O GLU A 152	29.980	8.096 123.683	1.00 31.23
*** ***		•			-

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```
ATOM
          1189
                N
                    TYR A 153
                                     32.054
                                               7.253 123.583
                                                               1.00 28.72
  MOTA
          1190
                CA
                    TYR A 153
                                     32.066
                                               6.805 124.971
                                                               1.00 31.35
  ATOM
          1191
                CB
                    TYR A 153
                                     33.427
                                               6.204 125.307
                                                               1.00 31.56
  MOTA
          1192
                CG
                    TYR A 153
                                     33.617
                                               5.839 126.759
                                                               1.00 33.17
                    TYR A 153
  MOTA
         1193
                CD1
                                     33.111
                                               4.647 127.280
                                                               1.00 35.43
  MOTA
          1194
                CE1
                    TYR A 153
                                     33.321
                                               4.298 128.619
                                                               1.00 33.52
         1195
  ATOM
                CD2
                    TYR A 153
                                     34.329
                                              6.677
                                                     127.611
                                                               1.00
                                                                    34.29
 MOTA
         1196
                CE2
                    TYR A 153
                                     34.544
                                              6.342 128.944
                                                               1.00 35.34
 ATOM
         1197
                CZ
                    TYR A 153
                                              5.154 129.444
                                     34.041
                                                               1.00 37.50
         1198
 MOTA
                    TYR A 153
TYR A 153
               OH
                                    34.260
                                              4.835
                                                    130.767
                                                               1.00
                                                                    30.10
 ATOM
         1199
               C
                                    31.828
                                              8.022 125.857
                                                               1.00 32.71
 ATOM
         1200
               0
                    TYR A 153
                                              7.988 126.787
                                    31.026
                                                               1.00 29.14
 MOTA
         1201
               N
                    LEU A 154
                                    32.538
                                              9.102 125.552
                                                               1.00
                                                                    29.65
 ATOM
         1202
               CA
                    LEU A 154
                                    32.413
                                             10.332 126.310
                                                               1.00 32.87
         1203
 MOTA
               CB
                   LEU A 154
                                    33.477
                                                    125.847
                                             11.329
                                                               1.00 31.46
 ATOM
         1204
               CG
                    LEU A 154
                                    34.910
                                             11.053
                                                    126.324
                                                               1.00
                                                                    29.68
 MOTA
         1205
               CD1
                   LEU A 154
                                    35.898
                                             11.953 125.605
                                                              1.00 29.29
 ATOM
         1206
                   LEU A 154
               CD2
                                    34.989
                                             11.278
                                                    127.829
                                                               1.00 27.19
         1207
 ATOM
               С
                                    31.020
                    LEU A 154
                                             10.952 126.232
                                                              1.00
                                                                    34.63
 ATOM
         1208
               0
                    LEU A 154
                                    30.475
                                             11.379 127.250
                                                              1.00 32.58
         1209
                   ARG A 155
 ATOM
               N
                                             10.999
                                    30.443
                                                    125.035
                                                              1.00 36.63
 ATOM
         1210
               CA
                   ARG A 155
                                    29.107
                                             11.569 124.869
                                                              1.00 38.36
 ATOM
         1211
               CB
                   ARG A 155
                                    28.661
                                             11.502 123.405
                                                              1.00 36.32
 ATOM
         1212
               CG
                   ARG A 155
                                    29.581
                                             12.253 122.460
                                                              1.00 43.15
 ATOM
         1213
               CD
                   ARG A 155
                                    29.100
                                             12.201 121.023
                                                              1.00 41.10
 ATOM
        1214
               NE
                   ARG A 155
                                    27.936
                                             13.047 120.768
                                                              1.00 44.00
 ATOM
        1215
               CZ
                   ARG A 155
                                    27.331
                                             13.140 119.583
                                                              1.00 54.07
        1216
 ATOM
               NH1
                   ARG A 155
                                    27.772
                                            12.441 118.540
                                                              1.00 51.61
        1217
 ATOM
               NH2
                   ARG A 155
                                    26.291
                                            13.948 119.424
                                                              1.00 51.76
 ATOM
        1218
                   ARG A 155
                                    28.112
                                            10.821 125.745
                                                              1.00 36.25
        1219
 MOTA
                   ARG A 155
                                    27.270
                                            11.433 126.397
                                                              1.00 39.00
 ATOM
        1220
              N
                   LYS A 156
                                    28..213
                                             9.496 125.765
                                                              1.00 36.48
 ATOM
        1221
              CA
                   LYS A 156
                                    27.315
                                             8.698 126.587
                                                              1.00 39.06
ATOM
        1222
              CB
                   LYS A 156
                                    27.460
                                             7:213 126.256
                                                              1.00 41.88
        1223
ATOM
              CG
                   LYS A 156
                                    26.672
                                             6.816 125.020
                                                              1.00 51.15
ATOM
        1224
              CD
                  LYS A 156
                                    27.169
                                             7.505 123.781
                                                              1.00 55.56
ATOM
        1225
              CE
                   LYS A 156
                                    26.117
                                             7.502 122.676
                                                              1.00 55.63
        1226
              NZ
MOTA
                  LYS A 156
                                   24.993
                                             8.425 123.013
                                                              1.00 49.15
ATOM
        1227
              С
                  LYS A 156
                                    27.527
                                             8.932 128.076
                                                             1.00 39.91
        1228
MOTA
              0
                  LYS A 156
                                   26.636
                                             8.658 128.876
                                                             1.00 37:01
MOTA
        1229
              N
                  LYS A 157
                                   28.703
                                             9.431 128.448
                                                             1.00 37.73
                  LYS A 157
MOTA
        1230
              CA
                                   28.985
                                             9.725 129.847
                                                             1.00 36.52
        1231
              CB
MOTA
                  LYS A 157
                                   30.493
                                             9.700 130.122
                                                             1.00 35.64
MOTA
        1232
              CG
                  LYS A 157
                                   31.094
                                             8.308 130.174
                                                             1.00 35.44
ATOM
        1233
              CD
                  LYS A 157
                                   30.509
                                             7.510 131.335
                                                             1.00 31.28
        1234
              CE
ATOM
                  LYS A 157
                                   31.077
                                             6.106 131.388
                                                             1.00 31.48
        1235
ATOM
             NZ
                  LYS A 157
                                   30.464
                                             5.310 132.493
                                                             1.00 36.39
ATOM
       1236
              С
                  LYS A 157
                                   28.423
                                            11.097 130.197
                                                             1.00 38.12
       1237
              0
MOTA
                  LYS A 157
                                   28.531
                                            11.547
                                                   131.336
                                                             1.00 37.61
       1238
MOTA
             N
                  GLY A 158
                                   27.842
                                           11.768 129.205
                                                             1.00 36.27
ATOM
       1239
             CA
                  GLY A 158
                                   27.257
                                           13.074 129.452
                                                             1.00 34.31
MOTA
       1240
             C
                  GLY A 158
                                           14.293 128.894
                                   27.972
                                                             1.00 36.36
ATOM
       1241
             0
                  GLY A 158
                                   27.438
                                           15.399 128.963
                                                             1.00
                                                                  32.96
MOTA
       1242
             N
                  PHE A 159
                                   29.170
                                           14.117 128.344
                                                             1.00 33.89
ATOM
       1243
             CA
                  PHE A 159
                                                  127.796
                                   29.892
                                           15.260
                                                             1.00 30.29
             CB
       1244
MOTA
                                           14.892 127.504
                  PHE A 159
                                  31.346
                                                             1.00 28.62
MOTA
       1245
             CG
                  PHE A 159
                                  32.137
                                           14.555 128.730
                                                             1.00 28.80
ATOM
       1246
             CD1
                 PHE A 159
                                  32.043
                                           13.300
                                                  129.310
                                                             1.00 30.41
       1247
MOTA
             CD2
                 PHE A 159
                                  32.951
                                           15.513 129.327
                                                            1.00
                                                                  29.37
ATOM
       1248
             CE1
                 PHE A 159
                                  32.749
                                           12.996 130.472
                                                            1.00 34.42
                 PHE A 159
       1249
             CE2
MOTA
                                  33.661
                                           15.223
                                                  130.488
                                                            1.00 31.10
       1250
                 PHE A 159
MOTA
             CZ
                                  33.561
                                           13.963
                                                  131.062
                                                                  32.32
                                                            1.00
MOTA
       1251
             С
                 PHE
                     A 159
                                  29.224
                                           15.786 126.536
                                                            1.00 28.88
       1252
                     A 159
                 PHE
             0
                                           15.003 125.705
ATOM
                                  28.765
                                                            1.00 27.71
       1253
                 LYS A 160
MOTA
             N
                                  29.180
                                                  126.402
                                           17.110
                                                            1.00
                                                                  30.20
       1254
ATOM
                     A 160
                                  28.550
                                           17.766 125.254
                                                            1.00 33.98
```

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Figure 18-20

ATO	M 1255 CB LYS A 160	27.390 18.653 125 719 1 00 36 8
ATO	M 1256 CG LYS A 160	1.00 36.8
ATO	100	
ATO		25.105 18.850 126.723 1.00 48.5
ATO:	M 1259 NZ LYS A 160	25.500 20.003 127.651 1.00 50.8
		25.924 19.534 129.008 1.00 49 70
ATO		29.484 18.616 124.394 1.00 32 50
ATO		29.093 19.085 123.327 1 00 30 9
ATO		30.700 18.846 124.867 1.00 31.4
ATOI	M 1263 CA ARG A.161	31.665 19.626 124.108 1.00 29.97
OTA	M 1264 CB ARG A 161	21.00 29.9
ATO		70'486
ATO		= 1.00 37.03
· ATON		21 100 39.01
ATON		31.158 23.503 126.341 1.00 43.76
ATON		30.389 23.337 127.414 1.00 43.33
		29.117 22.985 127.274 1.00 45 95
ATOM		30.893 23.518 128.627 1.00 43.74
ATOM		33.009 18.910 124.161 1.00 32 71
ATOM		33.792 19.090 125.092 7 00 29 53
ATOM	1 1273 N ILE A 162	33.257 18.087 123.149 1.00 32.50
ATOM		
MOTA		33.147 15.355 123.879 1.00 30.25
		32.564 13.977 123.635 1.00 34.27
MOTA		35.353 17.816 121.886 1.00 26.46
ATOM		34.876 17.973 120.762 1.00 27.88
MOTA		36.626 18.067 122.168 1.00 23 13
MOTA		37.575 18.534 121.156 1.00 25.74
ATOM		38.384 19.729 121.681 1.00 26.25
ATOM	1284 CG LEU A 163	39.626 20.138 120.862 1.00 24.39
MOTA	1285 CD1 LEU A 163	
ATOM	1286 CD2 LEU A 163	
MOTA	1287 C LEU A 163	40.361 21.252 121.560 1.00 27.76 38.547 17.416 120.792 1.00 27.09
ATOM	1288 O LEU A 163	
ATOM	1289 N TYR A 164	
ATOM		38.808 17.257 119.496 1.00 26.97
ATOM		39.747 16.241 119.010 1.00 26.97
		39.021 15.181 118.179 1.00 23.38
ATOM	1292 CG TYR A 164	39.944 14.146 117.565 1.00 21.76
MOTA	1293 CD1 TYR A 164	40.563 13.179 118.353 1.00 22.49
ATOM	1294 CE1 TYR A 164	41.419 12.224 117.794 1.00 22.90
ATOM	1295 CD2 TYR A 164	40.202 14.142 116.194 1.00 18.74
MOTA	1296 CE2 TYR A 164	41.060 13.190 115.616 1.00 23.36
ATOM	1297 CZ TYR A 164	
ATOM	1298 OH TYR A 164	2.00 21.00
MOTA	1299 C TYR A 164	
MOTA	1300 O TYR A 164	
ATOM	1301 N ILE A 165	40.473 .17.511 117.112 1.00 19.75
MOTA		42.057 16.843 118.551 1.00 25.61
		43.149 17.462 117.804 1.00 24.43
ATOM	1303 CB ILE A 165	43.963 18.396 118.717 1.00 26 41
MOTA	1304 CG2 ILE A 165	45.127 19.017 117.937 1.00 19.36
MOTA	1305 CG1 ILE A 165	43.035 19.482 119.274 1.00 23.36
ATOM	1306 CD1 ILE A 165	43.685 20.402 120.299 1.00 25.05
ATOM	1307 C ILE A 165	1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1
ATOM	1308 O ILE A 165	
ATOM	1309 N ASP A 166	
ATOM	1310 CA ASP A 166	12 200
ATOM	1311 CB ASP A 166	45.022 15.386 115.228 1.00 27.11
ATOM		44.140 14.765 114.137 1.00 28.56
		44.699 13.461 113.599 1.00 34.59
ATOM	1313 OD1 ASP A 166	45.831 13.456 113.068 1.00 30.37
ATOM	1314 OD2 ASP A 166	43.995 12.437 113.717 1.00 23.27
ATOM	1315 C ASP A 166	46.319 15.924 114.614 1.00 24.47
ATOM	1316 O ASP A 166	1,00 24.47
ATOM	1317 N LEU A 167	45 450
ATOM	1318 CA LEU A 167	1,00 25.45
ATOM	1319 CB LEU A 167	500
ATOM		49.682 16.382 115.887 1.00 21.90
MION	1320 CG LEU A 167	49.143 17.444 116.858 1.00 26.62

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Figure 18-21

MOTA	1321 CD1 LEU A 167	50.249	7 17.845 117.821	1.00 25.88
ATOM		48.658		
ATOM	- · - · ·			
		49.405		
MOTA		50.504		
MOTA		48.736	-	
ATOM-		49.244	12.975 112 <i>.</i> 555	1.00 24.59
ATOM	1327 CB ASP A 168	48.209	11.852 112.410	
ATOM	1328 CG ASP A 168	48.722		
MOTA	1329 C ASP A 168	49,423		
ATOM	1330 O ASP A 168	48.629		
ATOM	1331 OD1 ASP A 168	49.085		
ATOM	1332 OD2 ASP A 168			
		48.777		
ATOM		50.448		
ATOM	1334 CA ALA A 169	50.693		
ATOM	1335 CB ALA A 169	52.068	13.498 108.601	1.00 21.17
MOTA	1336 C ALA A 169	49.612	13.636 108.093	1.00 26.57
ATOM	1337 O ALA A 169	49.641	14.204 107.000	1.00 26.90
ATOM	1338 N HIS A 170	48.673	12.746 108.406	1.00 21.63
MOTA	1339 CA HIS A 170	47.592	12.445 107.468	1.00 24.79
ATOM	1340 C HIS A 170	46.243	12.867 108.045	1.00 20.98
ATOM	1341 O HIS A 170	46.044	12.849 109.255	
ATOM	1342 CB HIS A 170	47.550		1.00 24.12
			10.950 107.131	1.00 23.17
MOTA	1343 CG HIS A 170	48.830	10.420 106.570	1.00 30.28
MOTA	1344 ND1 HIS A 170	49.842	9.982 107.385	1.00 31.00
MOTA	1345 CE1 HIS A 170	50.825	9.634 106.577	1.00 24.33
ATOM	1346 CD2 HIS A 170	49.224	10.329 105.273	1.00 22.88
ATOM	1347 NE2 HIS A 170	50.502	9.828 105.285	1.00 21.89
MOTA	1348 N HIS A 171	45.317	13.231 107.171	1.00 21.14
ATOM	1349 CA HIS A 171	43.993	13.661 107.591	1.00 25.57
MOTA	1350 CB HIS A 171	43.234	14.242 106.404	1.00 22.47
ATOM	1351 CG HIS A 171	41.857	14.719 106.746	1.00 29.75
ATOM	1352 CD2 HIS A 171	41.433	15.648 107.634	1.00 25.58
MOTA	1353 ND1 HIS A 171	40.721	14.201 106.160	
ATOM	1354 CE1 HIS A 171 .			1.00 28.90
		39.656	14.787 106.676	1.00 25.35
ATOM		40.060	15.669 107.573	1.00 32.40
ATOM	1356 C HIS A 171	43.169	12.533 108.204	1.00 29.61
ATOM	1357 O HIS A 171	43.169	11.411 107.698	1.00 27.62
ATOM	1358 N CYS A 172	42.461	12.852 109.286	1.00 26.52
ATOM	1359 CA CYS A 172	41.610	11.897 109.987	1.00 24.82
MOTA	1360 CB CYS A 172	41.460	12.322 111.456	1.00 29.47
ATOM	1361 SG CYS A 172	40.959	14.065 111.717	1.00 25.69
ATOM	1362 C CYS A 172	40.237	11.797 109.314	1.00 28.21
ATOM	1363 O CYS A 172	39.211	12.131 109.914	1.00 26.78
ATOM	1364 N ASP A 173	40.213	11.332 108.066	1.00 22.05
ATOM	1365 CA ASP A 173	38.949	11.217 107.350	1.00 27.39
	1366 CB ASP A 173			
ATOM		39.167	10.646 105.931	1.00 30.47
MOTA		39.824	9.264 105.922	1.00 29.77
MOTA	1368 OD1 ASP A 173	39.886	8.658 104.830	1.00 21.14
MOTA	1369 OD2 ASP A 173	40.288	8.787 106.978	1.00 30.04
ATOM	1370 C ASP A 173	37.895	10.400 108.105	1.00 27.86
MOTA	1371 O ASP A 173	36.720	10.762 108.120	1.00 23.47
ATOM	1372 N GLY A 174	38.309	.9.315 108.753	1.00 25.84
ATOM	1373 CA GLY A 174	37.344	8.513 109.490	1.00 28.49
ATOM	1374 C GLY A 174	36.694	9.296 110.619	1.00 26.14
ATOM	1375 O GLY A 174	35.475	9.287 110.780	
	1376 N VAL A 175			1.00 21.39
ATOM		37.510	9.984 111.409	1.00 27.24
MOTA	1377 CA VAL A 175	36.995	10.773 112.523	1.00 25.53
MOTA	1378 CB VAL A 175	38.137	11.299 113.401	1.00 30.54
MOTA	1379 CG1 VAL A 175	37.565	12.105 114.566	1.00 28.02
ATOM	1380 CG2 VAL A 175	38.973	10.129 113.911	1.00 21.30
ATOM	1381 C VAL A 175	36.163	11.955 112.035	1.00 25.01
ATOM	1382 O VAL A 175	35.130	12.282 112.623	1.00 21.60
ATOM	1383 N GLN A 176	36.601	12.594 110.957	1.00 25.43
	1384 CA GLN A 176	35.854	13.730 110.426	
ATOM		36.554		1.00 26.12
ATOM			14.336 109.205	1.00 24.71
MOTA	1386 CG GLN A 176	35.682	15.349 108.469	1.00 26.68

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Figure 18-22

MOTA	1387 CD GLN	A 176	36.385	16.00	2 107.306	1.00 29.54
MOTA	1388 OE1 GLN	A 176	37.382		4 107.486	
MOTA	1389 NE2 GLN	A 176	35.872		6 106.099	
ATOM	1390 C GLN	A 176	34.446		6 110.029	
ATOM	1391 O GLN	A 176	33.481		1 110.319	1.00 25.93
MOTA		A 177	34.330		3 109.362	1.00 32.21
ATOM		A 177	33.027		108.915	1.00 32.72
MOTA		A 177	33.181		108.053	1.00 34.20
MOTA		A 177	31.905		107.329	1.00 39.40
MOTA		A 177	32.060		106.497	1.00 33.40
ATOM	1397 OE1 GLU		32.056		100.437	1.00 41.42
ATOM		A 177	32.206		107.073	
ATOM		A 177	32.128		110.099	1.00 42.35
ATOM		A 177	. 30.945		110.093	1.00 30.54
ATOM		A 178	32.707		110.093	1.00 25.39
ATOM		A 178	31.971			1.00 27.03
ATOM		A 178	32.905		112.303	1.00 30.67
ATOM		A 178		3.036	113.289	1.00 30.49
			31.261		113.003	1.00 33.21
ATOM		A 178	30.145		113.493	1.00 28.64
MOTA		A 179	31.888		113.055	1.00 29.27
MOTA		A 179	31.256		113.751	1.00 30.49
ATOM		A 179	32.071		115.001	1.00 24.44
ATOM		A 179	32.469		115.781	1.00 24.43
MOTA	1410 CD1 PHE		33.749		115.657	1.00 25.04
ATOM	1411 CD2 PHE 2		31.536		116.563	1.00 23.09
MOTA		A 179	34.103		116.293	1.00 16.56
ATOM		A 179	31.881		117.204	1.00 26.38
MOTA		A 179	33.170		117.067	1.00 20.30
ATOM		A 179	31.079		112.891	1.00 31.00
ATOM		179	31.006		113.399	1.00 31.32
MOTA		180	30.980		111.584	1.00 31.68
ATOM	1418 CA TYR A		30.829		110.646	1.00 32.76
ATOM	1419 CB TYR 7		30.931		109.213	1.00 35.12
MOTA	1420 CG TYR A		31.331		108.172	1.00 36.27
ATOM	1421 CD1 TYR A		30.427		107.204	1.00 37.31
ATOM	1422 CE1 TYR A		30.801		106.244	1.00 34.81
MOTA	1423 CD2 TYR A		32.624		108.154	1.00 36.32
MOTA	1424 CE2 TYR A		33.007		107.203	1.00 37.83
ATOM	1425 CZ TYR A		32.088		106.250	1.00 36.05
ATOM	1426 OH TYR A		32.446		105.323	1.00 28.04
ATOM	1427 C TYR A		29.518	16.696	110.825	1.00 30.94
ATOM	1428 O TYR A		29.459		110.560	1.00 30.42
ATOM	1429 N ASP A		28.473		111.299	1.00 31.56
ATOM	1430 CA ASP A		27.180		111.444	1.00 37.30
ATOM	1431 CB ASP A		26.086	15.833	110.807	1.00 37.68
ATOM	1432 CG ASP A		25.645	14.689	111.705	1.00 39.86
ATOM	1433 OD1 ASP A		26.505	13.963	112.233	1.00 43.25
ATOM	1434 OD2 ASP A		24.425		111.871	1.00 46.56
ATOM	1435 C ASP A		26.754		112.866	1.00 36.81
ATOM	1436 O ASP A		25.571	17.286	113.109	1.00 33.91
ATOM	1437 N THR A		27.689	17.066	113.810	1.00 40.86
ATOM	1438 CA THR A	182	27.327		115.184	1.00 38.27
ATOM	1439 CB THR A		27.433		116.133	1.00 37.99
MOTA	1440 OG1 THR A		27.013		117.448	1.00 35.64
ATOM	1441 CG2 THR A	182	28.869	15.684	116.194	1.00 35.61
ATOM	1442 C THR A		28.177	18.546	115.746	1.00 39.51
ATOM	1443 O THR A	182	29.365	18.673	115.433	1.00 40.07
ATOM	1444 M ASP A		27.557		116.582	1.00 37.01
ATOM	1445 CA ASP A		28.250		117.181	1.00 37.74
ATOM	1446 CB ASP A		27.313		117.228	1.00 35.56
ATOM	1447 CG ASP A		26.136	21.493		1.00 38.01
ATOM	1448 OD1 ASP A		25.614		118.210	1.00 34.94
ATOM	1449 OD2 ASP A		25.720	22.470		1.00 34.34
ATCM ·	1450 C ASP A		28.762	20.161		1.00 35.27
ATCM	1451 O ASP A		29.337	21.015		1.00 35.27
ATOM	1452 N GLN A		28.562	18.917		1.00 35.10
A. C.	GM. h		20.502	-U.J.		1.00 33.10

ATOM	1453	ÇA	GLN A	184	29.030		120.333	1.00 35.16
MOTA	1454	CB	GLN A	184	28.155	17.382	120.906	1.00 36.94
ATOM	1455	CG	GLN A		26.663	17.718	120.988	1.00 38.34
	1456	CD	GLN A		25.881		121.838	1.00 43.68
MOTA							121.696	1.00 35.48
MOTA	1457	OE1	GLN A		26.027			1.00 51.06
ATOM	1458	NE2	GLN A		25.036		122.723	
ATOM	1459	С	GLN A	184	30.479		120.253	1.00 36.32
ATOM	1460	0	GLN A	184	31.135	17.825	121.275	1.00 34.24
ATOM	1461	N	VAL A		30.976	17.883	119.028	1.00 34.51
	1462	CA	VAL A		32.348		118.804	1.00 33.59
ATOM		CB	VAL A		32.393		118.259	1.00 35.11
ATOM	1463				33.834		118.003	1.00 23.80
MOTA	1464	CG1	VAL A				119.242	1.00 26.00
MOTA	1465	CG2	VAL A		31.731	15.045	117.242	
MOTA	1466	С	VAL A	185	33.053		117.803	1.00 33.11
ATOM	1467	0	VAL A	185	32.545	18.593	116.714	1.00 27.73
ATOM	1468	N	PHE A	186	34.215	18.872	118.184	1.00 31.49
MOTA	1469	CA	PHE A		34.985	19.729	117.291	1.00 30.63
	1470	CB	PHE A		35.420	21.023	117.991	1.00 30.34
ATOM		CG	PHE A		36.008		117.051	1.00 30.22
ATOM	1471				35.265		116.656	1.00 32.23
MOTA	1472		PHE A			21 070	116.524	1.00 29.37
MOTA	1473	CD2	PHE A		37.284	21.079	110.324	
MOTA	1474	CE1	PHE A		35.785		115.748	1.00 27.87
ATOM	1475	CE2	PHE A	186	37.813		115.615	1.00 28.54
ATOM	1476	CZ	PHE A	186	37.064		115.227	1.00 30.80
ATOM	1477	С	PHE A	186	36.232		116.879	1.00 33.38
ATOM	1478	ō.	PHE A		36.952	18.426	117.729	1.00 28.30
	1479	N	VAL A		36.478		115.574	1.00 32.00
MOTA		CA	VAL A		37.645		115.060	1.00 29.70
MOTA	1480				37.252		114.019	1.00 30.03
MOTA	1481	CB	VAL A		38.510	16 400	113.405	1.00 27.77
ATOM	1482	CG1	VAL A					1.00 25.98
MOTA	1483	CG2	VAL A		36.410		114.672	
MOTA	1484	С	VAL A		38.604		114.392	1.00 31.03
ATOM	1485	0	VAL A	187	38.215		113.491	1.00 31.88
MOTA	1486	N	LEU A	188	39.850	19:157	114.857	1.00 24.88
ATOM	1487	CA	LEU A		40.899	20.010	114.304	1.00 26.92
	1488	CB	LEU A		41.468		115.361	1.00 27.04
ATOM	1489	CG	LEU A		42.823		114.963	1.00 25.15
MOTA			LEU A		42.686	22 315	113.648	1.00 18.53
ATOM	1490	CD1			43.330	22.315	116.068	1.00 28.81
ATOM	1491	CD2	LEU A				113.815	1.00 31.22
ATOM	1492	С	LEU A		42.022			1.00 25.83
ATOM	1493	0	LEU A		42.579		114.587	
MOTA	1494	N	SER A	189	42.369	19.230	112.540	1.00 30.53
ATOM	1495	CA	SER A	189	43.429	18.399	112.007	1.00 30.13
MOTA	1496	CB	SER A	18 <i>9</i>	42.821	17.249	111.199	1.00 33.41
ATOM	1497	OG	SER A	189	43.837	16.474	110.588	1.00 32.98
	1498	c	SER A		44.448	19.120	111.143	1.00 27.94
MOTA	1499	ō	SER A		44.084	19.891	11).253	1.00 22.14
ATOM		N	LEU A		45.728	18.877	11 . 423	1.00 24.80
ATOM	1500				46.805	10.077	110.614	1.00 22.23
MOTA	1501	CA	LEU A				111.459	1.00 23.69
MOTA		· CB	LEU A		47.955			1.00 28.92
MOTA.	1503	CG	LEU A		47.733	21.075	112.522	
ATOM	1504		LEU A		49.070		112.740	1.00 23.01
ATOM	1505	CD2	LEU A	190	46.691	22.093	112.087	1.00 28.11
ATOM	1506	С	LEU A	190	47.300	18.210	109.872	1.00 22.69
	1507	ō	LEU A	190	47.416	17.141	110.465	1.00 16.55
MOTA	1508	N	HIS A		47.599	18.353	108.587	1.00 19.22
ATOM			HIS A		48.046	17.210		1.00 23.28
MOTA	1509	CA	MID A	101	46.870	16.242		1.00 15.58
ATOM	1510	CB	HIS A	171				1.00 24.16
ATOM	1511	CG	HIS A	191	45.591	16.915		1.00 17.71
ATOM	1512	CD2	HIS A	191	45.034	17.124	106.038	1.00 17.71
ATOM	1513	ND1	HIS A	191	44.695	17.419	108.176	1.00 23.76
ATOM	1514	CE1.	HIS A	191	43.644	17.913	107.545	1.00 19.78
ATOM	1515	NE2	HIS A	191	43.823	17.746	106.246	1.00 27.87
	1516	C	HIS A	191	48.570	17.620	106.434	1.00 23.65
ATOM	1517	ò	HIS A	191	48.419	18.761	106.017	1.00 23.89
ATOM			GLN A	192	49.209	16.681		1.00 23.49
ATOM	1518	Ħ	GTW W	~~~				

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1.00 20.55
                                    49.718
                                            16.950 104.412
                   GLN A 192
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              CA
ATOM
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                                                              1.00 23.63
                   GLN A 192
                                    50.474
        1520
              CB
MOTA
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                                    51.528
                   GLN A 192
        1521
               CG
MOTA
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                                                              1.00 26.90
                                    52.110
                   GLN A 192
        1522
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MOTA
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                                                              1.00 20.21
                                    52.986
                  GLN A 192
        1523
              OE1
MOTA
                                                              1.00 23.52
                                    51.605
                                            12.765 104.828
                   GLN A 192
        1524
              NE<sub>2</sub>
MOTA
                                    48.478
                                            17.174 103.570
                                                              1.00 21.41
                   GLN A 192
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               C
MOTA
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                                                              1.00 20.15
                                    47.478
                   GLN A 192
ATOM
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                                                              1.00 24.36
                                    48.528
                                            18.167
                                                    102.692
                   SER A 193
        1527
              N
MOTA
                                            18.448 101.821
                                                              1.00 23.98
                                    47.397
                   SER A 193
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              CA
MOTA
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                                                              1.00 24.60
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MOTA
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              CB
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                                                    99.861
                   SER A 193
        1530
              OG
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                                            17.200 101.045
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ATOM
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                   PRO A 194
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MOTA
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                                    43.554
                   PRO A 194
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               CG
MOTA
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                                                              1.00 30.75
                                            15.825
                   PRO A 194
                                    45.527
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MOTA
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                                                                   30.04
                                    45.420
                                            14.830
                                                     98.041
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MOTA
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                                            16.991
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MOTA
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                                                              1.00 31.11
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MOTA
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                                                     96.571
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MOTA
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                                                     96.352
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MOTA
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                                                     96.434
                                    47.139
                                             21.425
                   GLU A 195
MOTA
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                                             21.605
                                                     95.825
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        1546
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                   GLU A 195
MOTA
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                                                     96.552
                                                              1.00 30.81
                                             16.193
                   GLU A 195
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MOTA
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                                                              1.00 37.17
                                                     95.426
                   GLU A 195
                                    47.582
                                             15.705
ATOM
        1548
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                                                              1.00 25.01
                                                     97.515
                                    48.377
                                             15.922
                   TYR A 196
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MOTA
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                                             15.053
                                                     97.242
                                    49.517
                   TYR A 196
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MOTA
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                                                     97.223
                                    50.810
                   TYR A 196
MOTA
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               CB
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                                    51.255
                                             16.424
                   TYR A 196
MOTA
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                                                     99.476
                                    51.957
                                             15.625
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 MOTA
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                                             16.110 100.734
                                    52.338
                   TYR A 196
               CEI
        1554
 ATOM
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                                             17.731
                                                     98.958
                                    50.944
                   TYR A 196
        1555
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 MOTA
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                                                              1.00 25.95
                                    51.320
        1556
                   TYR A 196
               CE2
 ATOM
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                                             17.409 101.096
                                    52.012
                   TYR A 196
               CZ
 MOTA
        1557
                                             17.879
                                                    102.345
                                                               1.00 25.50
                                    52.356
                   TYR A 196
        1558
               OH
 MOTA
                                                              1.00 27.05
                                             13.906
                                                     98.229
                                    49.670
                   TYR A 196
        1559
               С
 ATOM
                                                              1.00 24.02
                                                     98.096
                                    50.585
                                             13.088
                   TYR A 196
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        1560
 MOTA
                                                     99.214
                                                              1.00 22.10
                                             13.822
                   ALA A 197
                                    48.785
        1561
               N
 MOTA
                                             12.760 100.199
                                                              1.00 24.90
                                    48.928
                   ALA A 197
        1562
               CA
 MOTA
                                                              1.00 27.83
                                    49.627
                                             13.307 101.437
                   ALA A 197
        1563
               CB
 MOTA
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                                                    100.608
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                   ALA A 197
 ATOM
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                                                               1.00 22.82
                                    46.553
                   ALA A 197
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 ATOM
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                                             10.849 101.102
                                    47.795
                   PHE A 198
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        1536
 ATOM
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                                    46.663
                                             10.072 101.580
        1567
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               CA
 MOTA
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                                                               1.00 30.66
                   PHE A 198
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        1568
               CB
 MOTA
                                              7.766 102.399
                                                               1.00 29.61
                                    46.009
                   PHE A 198
               CG
 MOTA
         1569
                                                               1.00 28.76
                                    45.496
                                              6.879
                                                    101.463
                   PHE A 198
        1570
               CD1
 ATOM
                                              7.822 103.657
                                                               1.00 28.43
                                    45.426
                   PHE A 198
               CD2
         1571
 MOTA
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                                                               1.00 35.72
                                    44.415
                       A 198
        1572
               CE1
                   PHE
 MOTA
                                                               1.00 34.62
                                    44.340
                                              7.004 103.970
                   PHE A 198
        1573
               CE2
 MOTA
                                              6.121 103.029
                                                               1.00 35.73
                                    43.837
                   PHE A 198
         1574
               CZ
 MOTA
                                             10.814 102.802
                                                               1.00 28.95
                       A 198
                                    46.121
                   PHE
        1575
               С
 MOTA
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                                    46.892
                                             11.347
                                                     103.596
                   PHE A 198
        1576
               a
 MOTA
                                             10.905 102.941
                                                               1.00 28.27
                                    44.792
                   PRO A 199
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 ATOM
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                                                               1.00 33.97
                   PRO A 199
                                    44.100
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               CD
 ATOM
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                                    43.813
                                             10.364 102.008
                   PRO A 199
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               CA
 ATOM
                                             10.312 102.858
                                                               1.00 29.84
                                    42.550
                   PRO A 199
               CB
        1580
 MOTA
                                                               1.00
                                                                    37.32
                   PRO A 199
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                                             11.612 103.592
               CG
        1581
 ATOM
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                                                               1.00 35.02
                                    43.773
                   PRO A 199
        1582
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 MOTA
                                             12.631 101.280
                                                               1.00 48.84
                   PRO A 199
                                    44.052
               0
        1583
 MOTA
                                                      99.734
                                                               1.00 33.64
                          200
                                    43.441
                                             11.156
                   PHE A
         1584
               N
 ATOM
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MOTA	1585	CA	PHE A 200		43.418	12.179	98.718	1.00 28.12
MOTA	1586	CB	PHE A 200		43.927	11.579	97.411	1.00 26.69
ATOM	1587				45.226	10.833	97.561	1.00 27.33
ATOM	1588				45.239	9.510	97.995	1.00 29.79
MOTA	1589				46.439	11.461	97.302	1.00 24.38
ATOM	1590	CE:	1 PHE A 200		46.444	8.820	98.168	1.00 29.45
ATOM	1591	CE:	2 PHE A 200		47.651	10.782	97.473	1.00 31.41
ATOM	1592	CZ	PHE A 200		47.653	9.458	97.906	1.00 29.64
	1593	c	PHE A 200		42.042	12.795	98.518	1.00 26.15
MOTA								
MOTA	1594	0	- PHE A 200		41.935	13.889	97.986	1.00 27.96
ATOM	1595	N	GLU A 201		41.002	12.101	98.979	1.00 28.52
ATOM	1596	CA	GLU A 201		39.614	12.534	98.806	1.00 35.04
ATOM	1597	CB	GLU A 201	•	38.695	11.316	98.810	1.00 33.61
ATOM	1598	CG	GLU A 201		39.087	10.240	97.838	1.00 37.80
ATOM	1599	CD	GLU A 201		38.222	9.016	97.997	1.00 43.48
ATOM	1600	OE:			36.992	9.142	97.825	1.00 40.96
	1601	OE						1.00 44.17
MOTA					38.772	7.937	98.298	
MOTA	1602	C	GLU A 201		39.077	13.516	99.837	1.00 36.30
MOTA	1603	0	GLU A 201		38.087	14.206	99.592	1.00 36.47
MOTA	1604	N	LYS A 202		39.693	13.552	101.007	1.00 34.63
MOTA	1605	CA	LYS A 202		39.229	14.460	102.030	1.00 34.09
ATOM	1606	CB	LYS A 202		38.294	13.729	102.992	1.00 40.88
ATOM	1607	CG	LYS A 202		37.011		102.292	1.00 43.17
	1608	CD	LYS A 202		35.935		103.230	1.00 47.39
ATOM			LYS A 202		34.628	12.663	102.469	1.00 47.74
ATOM	1609	CE						
MOTA	1610	NZ	LYS A 202		33.504		103.378	1.00 53.56
MOTA	1611	С	LYS A 202		40.382		102.753	1.00 36.27
MOTA	1612	0	LYS A 202		41.520		102.613	1.00 28.06
ATOM	1613	N	GLY A 203		40.080	16.152	103.509	1.00 31.91
MOTA	1614	CA	GLY A 203		41.115	16.862	104.228	1.00 33.75
ATOM	1615	С	GLY A 203	•	41.288	18.288	103.729	i.00 30.54
ATOM	1616	0	GLY A 203		42.174		104.200	1.00 28.04
ATOM	1617	N	PHE A 204		40.458	18.713		1.00 29.93
		CA	PHE A 204		40.557	20.077		1.00 35.76
ATOM	1618							
MOTA	1619	CB	PHE A 204		39.863	20.217		1.00 31.41
MOTA	1620	CG	PHE A 204		40.498	19.416	99.803	1.00 31.06
MOTA	1621	CD1			40.169	18.075	99.618	1.00 35.66
ATOM	1622	CD2	PHE A 204		41.431	20.002	98.955	1.00 30.79
MOTA	1623	CE1	PHE A 204		40.761	17.329	98.597	1.00 35.20
ATOM	1624	CE2	PHE A 204		42.033	19.267	97.931	1.00 36.08
ATOM	1625	CZ	PHE A 204		41.697	17.928	97.751	1.00 36.54
ATOM	1626	c	PHE A 204		39.967	21.103	103.231	1.00 37.30
	1627	ō	PHE A 204		39.088	20.786	104.040	1.00 33.56
MOTA	1628	N	LEU A 205		40.451	22.337	103.128	1.00 38.52
MOTA					40.012		103.120	1.00 36.81
ATOM	1629	CA	LEU A 205			23.427		
ATOM	1630	CB	LEU A 205		40.801	24.695	103.659	1.00 34.73
MOTA	1631	CG	LEU A 205		40.496		104.479	1.00 40.98
MOTA	1632		LEU A 205		-40.690		105.965	1. <u>00</u> 39.87
ATOM	1633	CD2	LEU A 205		41.415	27.079	104.032	1.00 39.94
ATOM	1634	С	LEU A 205		38.520	23.728	103.925	1.00 36.58
MOTA	1635	0	LEU A 205		37.931	24.178	104.905	1.00 40.98
ATOM	1636	N	GLU A 206		37.909		102.774	1.00 36.07
	1637	CA	GLU A 206		36.486		102.586	1.00 36.30
MOTA					36.107		101.105	1.00 30.38
ATOM	1638	CB	GLU A 206					
ATOM	1639	CG	GLU A 206		36.890	24.473	100.131	1.00 48.04
MOTE	1640	CD	GLU A 206		38.307	23.980	99.868	1.00 51.87
MOTA	1641	0E1	GLU A 206		39.146	23.993	100.792	1.00 50.32
ATOM	1642	OE2	GLU A 206		38.581	23.569	98.716	1.00 56.69
ATOM	1643	С	GLU A 206		35.572	22.852	103.427	1.00 33.85
ATOM	1644	5	GLU A 206		34.433	23.213	103.718	1.00 26.22
	1645	И	GLU A 207		36.071		103.805	1.00 31.68
ATOM	1646	CA	GLU A 207		35.297		104.599	1.00 31.65
ATOM					36.000		104.566	1.00 34.15
ATOM	1647	CB	GLU A 207			19.307	102.700	
ATOM	1648	CG	GLU A 207		36.044		103.179	1.00 33.80
ATOM	1649	CD	GLU A 207		37.182		103.022	1.00 33.85
ATOM	1650	OE1	GLU A 207		37.487	17.025	103.995	1.00 33.22

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MOTA	1651 OE2 GLU A 207	37.760	17.688 101.916	1.00 35.48
MOTA	1652 C GLU A.207	35.182		
ATOM	1653 O GLU A 207	36.009	_	
MOTA		34.150		
MOTA	1655 CA ILE A 208	33.968		
MOTA	1656 CB ILE A 208	33.737	24.134 107.529	1.00 42.74
ATOM	1657 CG2 ILE A 208	33.717	24.762 108.914	1.00 48.29
MOTA	1658 CG1 ILE A 208	34.841		1.00 40.34
ATOM	1659 CD1 ILE A 208	36.207		
				1.00 46.23
MOTA		32.821		1.00 38.32
ATOM	1661 O ILE A 208	- 32.558		1.00 40.08
MOTA	1662 N GLY A 209	32.142	20.997 107.901	1.00 34.36
ATOM	1663 CA GLY A 209	31.047	20.374 108.620	1.00 33.32
ATOM	1664 C GLY A 209	29.699		1.00 37.87
MOTA	1665 O GLY A 209	29.579		1.00 40.56
ATOM	1666 N GLU A 210			
	·	28.676		1.00 37.38
MOTA	1667 CA GLU A 210	27.337		1.00 42.34
MOTA	1668 CB GLU A 210	27.008	19.012 106.823	1.00 42.73
MOTA	1669 CG GLU A 210	26.860	17.636 107.460	1.00 47.38
ATOM	1670 CD GLU A 210	26.633	16.532 106.443	1.00 52.68
ATOM	1671 OE1 GLU A 210	26.385	15.379 106.860	1.00 50.59
	1672 OE2 GLU A 210			
MOTA		26.711	16.810 105.226	1.00 53.78
MOTA	1673 C GLU A 210	26.287	20.114 108.938	1.00 42.90
ATOM	1674 O GLU A 210	26.516	19.577 110.022	1.00 45.94
ATOM	1675 N GLY A 211	25.130	20.702 108.654	1.00 43.16
ATOM	1676 CA GLY A 211	24.068	20.751 109.642	1.00 43.98
ATOM	1677 C GLY A 211	24.514	21.450 110.911	1.00 45.01
ATOM	1678 O GLY A 211	25.186	22.479 110.858	1.00 48.15
MOTA	1679 N LYS A 212		20.896 112.059	
		24.145		1.00 41.63
MOTA	1680 CA LYS A 212	24.528	21.495 113.328	1.00 45.07
ATOM	1681 CB LYS·A 212	23.913	20.715 114.490	1.00 46.59
ATOM	1682 CG LYS A 212	22.386	20.591 114.462	1.00 55.31
MOTA	1683 CD LYS A 212	21.651	21.945 114.481	1.00 57.42
ATOM .	1684 CE LYS A 212 .	21.749	22.696 113.151	1.00 59.71
ATOM	1685 NZ LYS A 212	21.051	24.017 113.178	
				1.00 57.43
MOTA		26.046	21.513 113.469	1.00 42.08
ATOM	1687 O LYS A 212	26.598	22.326 114.207	1.00 40.03
MOTA	1688 N GLY A 213	26.713	20.615 112.751	1.00 39.51
MOTA	1689 CA GLY A 213	28.163	20.538 112.817	1.00 40.11
ATOM	1690 C GLY A 213	28.888	21.519 111.916	1.00 38.25
ATOM	1691 O GLY A 213	30.122	21.575 111.913	1.00 34.70
ATOM	1692 N LYS A 214	28.131	22.295 111.143	1.00 37.31
MOTA	1693 CA LYS A 214	28.736		
			23.274 110.250	1.00 39.58
MOTA	1694 CB LYS A 214	27.656	24.017 109.463	1.00 44.69
MOTA	1695 CG LYS A 214	28.189	25.030 108.461	1.00 44.53
MOTA	1696 CD LY: A 214	27.047	25.704 107.720	1.00 47.71
ATOM	1697 CE LY. A 214	27.553	26.759 106.754	1.00 52.94
ATOM	1698 NZ LYC A 214	28.453	26.183 105.717	1.00 57.45
MOTA	1699 C LYS A 214	29.547	24.259 111.085	1.00 40.16
ATOM	1700 O LYS A 214	29.002	24.963 111.933	
MOTA	1701 N GLY A 215	30.851	24.295 110.846	1.00 36.57
MOTA	1702 CA GLY A 215	31.716	25.183 111.593	1.00 35.03
MOTA	1703 C GLY A 215	32.431	24.448 112.709	1.00 34.57
ATOM	1704 O GLY A 215	33.216	25.039 113.454	1.00 33.76
ATOM	1705 N TYR A 216	32.168	23.153 112.837	1.00 34.61
	1706 CA TYR A 216	32.816	22.378 113.885	1.00 35.00
MOTA				
MOTA	1707 CB TYR A 216	31.763	21.683 114.753	1.00 36.19
ATCM	1708 CG TYR A 216.	30.928	22.671 115.547	1.00 36.68
MOTA	1709 CD1 TYR A 216	29.961	23.462 114.925	1.00 35.83
ATOM	1710 CE1 TYR A 216	29.249	24.431 115.641	1.00 40.89
ATOM	1711 CD2 TYR A 216	31.163	22.869 116.910	1.00 41.50
	1712 CE2 TYR A 216	30.459	23.834 117.634	1.00 40.69
ATOM				
MOTA		29.505	24.612 116.994	1.00 40.17
ATOM	1714 OH TYR A 216	28.816	25.566 117.708	1.00 38.09
ATOM	1715 C TYR A 216	33.877	21.384 113.401	1.00 34.05
ATOM	1716 O TYR A 216	34.263	20.462 114.127	1.00 31.87

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1717
  ATOM
                 N
                     ASN A 217
                                      34.343
                                              21.580 112.170
                                                               1.00 29.90
  ATOM
          1718
                 CA
                     ASN A 217
                                     35.398
                                              20.748 111.606
                                                               1.00 30.02
          1719
  MOTA
                 CB
                     ASN A 217
                                     34.833
                                              19.727 110.615
                                                               1.00 26.46
  ATOM
          1720
                     ASN A 217
                                     35.897
                                              18.764 110.105
                                                               1.00
                                                                     30.13
          1721
  ATOM
                 OD1
                     ASN A 217
                                     36.558
                                              19.022 109.097
                                                               1.00 29.80
          1722
  MOTA
                 ND2
                     ASN A 217
                                     36.094
                                              17.659 110.831
                                                               1.00 19.92
  ATOM
          1723
                                     36.378
                     ASN A 217
                                              21.686 110.915
                                                               1.00 30.23
          1724
  MOTA
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                     ASN A 217
                                     35.983
                                              22.502 110.080
                                                               1.00 27.88
          1725
 MOTA
                N
                     LEU A 218
                                     37.655
                                              21.577 111.271
                                                               1.00 29.45
  ATOM
          1726
                 CA
                     LEU A 218
                                     38.670
                                              22.451 110.698
                                                               1.00 28.76
 MOTA
          1727
                CB
                     LEU A 218
                                     39.160
                                              23.444 111.753
                                                               1.00 29:02
 MOTA
          1728
                CG
                     LEU A 218
                                     39.513
                                              24.867 111.307
                                                               1.00 34.69
 ATOM
          1729
                CD1
                    LEU A 218
                                     40.432
                                              25.480
                                                     112.367
                                                               1.00 32.93
 ATOM
          1730
                CD2
                    LEU A 218
                                     40.197
                                              24.873 109.954
                                                               1.00 30.69
         1731
 MOTA
                C
                     LEU A 218
                                     39.870
                                              21.657 110.207
                                                               1.00 26.65
         1732
 ATOM
                     LEU A 218
                                     40.527
                                              20.981 110.999
                                                               1.00
                                                                    25.25
         1733
 MOTA
                N
                     ASN A 219
                                     40.151
                                              21.752 108.909
                                                               1.00 25.21
         1734
                CA
                                             21.069 108.294
 MOTA
                    ASN A 219
                                     41.287
                                                               1.00 21.91
         1735
 ATOM
                CB
                    ASN A 219
                                     40.875
                                             20.314 107.018
                                                               1.00 23.69
         1736
                    ASN A 219
 ATOM
                CG
                                    39.972
                                             19.144 107.298
                                                               1.00 27.88
         1737
                OD1
 ATOM
                                             18.440 108.289
                    ASN A 219
                                     40.153
                                                               1.00 29.28
         1738
 ATOM
                ND2
                    ASN A 219
                                    39.018
                                             18.900 106.407
                                                               1.00 24.48
         1739
 ATOM
                C
                    ASN A 219
                                    42.355
                                             22.074 107.906
                                                               1.00 23.46
         1740
 ATOM
                0
                    ASN A 219
                                    42.059
                                             23.073
                                                               1.00 28.17
                                                    107.259
         1741
 ATOM
                N
                    ILE A 220
                                    43.595
                                             21.804 108.287
                                                               1.00 23.90
         1742
                                    44.702
 MOTA
                CA
                    ILE A 220
                                             22.684 107.945
                                                               1.00 23.22
 ATOM
         1743
                CB
                   ILE A 220
                                    45.468
                                             23.131 109.212
                                                               1.00 28.73
         1744
               CG2 ILE A 220
 ATOM
                                    46.601
                                             24.078 108.831
                                                               1.00 26.01
         1745
 ATOM
               CG1
                    ILE A 220
                                    44.502
                                             23.783 110.212
                                                               1.00 26.36
         1746
 ATOM
               CD1
                    ILE A 220
                                    43.771
                                             25.004 109.688
                                                               1.00 25.74
 ATOM
         1747
               C
                    ILE A 220
                                    45.669
                                             21.929 107.018
                                                               1.00 25.29
         1748
                                             21.315 107.477
 ATOM
               0
                    ILE A 220
                                    46.631
                                                              1.00 20.44
         1749
 ATOM
               N
                    PRO A 221
                                    45.396
                                             21.924, 105.703
                                                               1.00 26.34
 ATOM
        1750
               CD
                    PRO A 221
                                    44.234
                                             22:497 104.999
                                                              1.00 28.22
        1751
                   PRO A 221
 ATOM
               CA
                                    46.271
                                             21.234 104.747
                                                              1.00 26.92
        1752
 ATOM
               CB
                   PRO A 221
                                    45.454
                                             21.279 103.457
                                                              1.00 27.81
ATOM
        1753
                   PRO A 221
                                    44.774
               CG
                                                              1.00 30.62
                                             22.622 103.582
        1754
ATOM
               С
                   PRO A 221
                                    47.595
                                            21.977
                                                    104.625
                                                              1.00 27.45
        1755
 MOTA
               0
                   PRO A 221
                                    47.603
                                            23.199 104.457
                                                              1.00 31.21
        1756
ATOM
               N
                   LEU
                       A 222
                                    48.704
                                            21.242 104.703
                                                              1.00 26.01
ATOM
        1757
               CA
                   LEU A 222
                                   50.038
                                            21.838 104.640
                                                              1.00 26.41
        1758
               CB
                   LEU A 222
MOTA
                                   50.726
                                            21.650 105.997
                                                              1.00 26.12
ATOM
        1759
               CG
                   LEU A 222
                                   49.960
                                            22.322 107.150
                                                              1.00 27.67
        1760
               CD1
                   LEU A 222
                                   50.531
MOTA
                                            21.899
                                                    108.497
                                                              1.00 30.97
ATOM
        1761
              CD2
                   LEU A 222
                                   50.024
                                            23.839 106.985
                                                              1.00 31.59
        1762
MOTA
                   LEU À 222
                                   50.911
                                            21.286 103.504
                                                              1.00 28.97
MOTA
        1763
              0
                   LEU A 222
                                   50.784
                                            20.128 103.117
                                                              1.00 27.95
                   PRO A 223
PRO A 223
        1764
ATOM
              N
                                   51.821
                                            22.116 102.964
                                                              1.00 31.52
        1765
              CD
MOTA
                                   52.059
                                            23.518 103.358
                                                              1.00 29.08
        1766
ATOM
              CA
                   PRO A 223
                                   52.727
                                            21.753 101.865
                                                              1.00 29.93
        1767
MOTA
              CB
                   PRO A 223
                                   53.265
                                            23.109 101.428
                                                              1.00 29.16
        1768
              CG
                  PRO A 223
MOTA
                                   53.458
                                                   102.779
                                                             .1.00 25.86
                                            23.771
MOTA
        1769
              С
                   PRO A 223
                                   53.862
                                            20.782 102.206
                                                             1.00 33.62
                  PRO A 223
        1770
MOTA
              0
                                   54.179
                                            20.531 103.376
                                                              1.00 26.55
       1771
MOTA
              N
                  LYS A 224
                                   54.479
                                            20.257
                                                    101.153
                                                             1.00 34.00
       1772
                                   55.595
ATOM
              CA
                  LYS A 224
                                            19.320
                                                   101.264
                                                             1.00 32.88
       1773
                  LYS A 224
              СB
MOTA
                                   55.938
                                            18.767
                                                     99.884
                                                             1.00 36.31
       1774
              CG
                  LYS
                                   54.761
ATOM
                      A 224
                                                    99.115
                                                             1.00 39.37
                                            18.204
       1775
                                                    97.658
              CD
                  LYS A 224
                                   55.150
MOTA
                                           17.998
                                                             1.00 45.23
       1776
                  LYS A 224
ATOM
              CE
                                   53.989
                                                             1.00 47.90
                                           17.478
                                                    96.835
       1777
                  LYS A 224
ATOM
             NZ
                                  54.331
                                           17.441
                                                    95.388
                                                             1.00 46.60
       1778
                  LYS A 224
                                  56.817
ATOM
                                           20.054
                                                   101.798
                                                             1.00 29.43
       1779
             0
                  LYS A 224
                                  56.933
MOTA
                                           21.270 101.640
                                                             1.00 24.10
       1780
                  GLY A 225
ATOM
             N
                                  57.735
                                           19.305 102.403
                                                             1.00 25.00
       1781
             CA
                  GLY A 225
ATOM
                                  58.947
                                           19.896 102.942
                                                             1.00 26.20
                  GLY A 225
       1782
ATOM
             C
                                  58.727
                                           20.792 104.154
                                                             1.00 29.44
```

ATOM	1783 O GLY A 225	59.610 21.562 104.528 1.00 29.09
ATOM		57.560 20.679 104.777 1.00 24.26
ATOM		57.212 21.488 105.951 1.00 25.39
MOTA	1786 CB LEU A 226	55.930 20.925 106.579 1.00 23.23
MOTA		55.172 21.757 107.611 1.00 28.28
MOTA	1788 CD1 LEU A 226	54.596 22.972 106.911 1.00 28.07
MOTA		54.036 20.933 108.226 1.00 27.49
ATOM		58.333 21.501 106.998 1.00 24.79
ATOM		58.902 20.450 107.299 1.00 26.15
ATOM		58.664 22.674 107.548 1.00 23.94
MOTA	1793 CA ASN A 227	59.702 22.733 108.578 1.00 24.89
MOTA		60.751 23.823 108.269 1.00 27.43
ATOM	1795 CG ASN A 227	60.190 25.231 108.334 1.00 31.62
MOTA	1796 OD1 ASN A 227	59.598 25.632 109.336 1.00 31.17
MOTA	1797 ND2 ASN A 227	60.395 25.998 107.267 1.00 28.06
MOTA	1798 C ASN A 227	59.076 22.957 109.960 1.00 23.16
ATOM	1799 O ASN A 227	57.873 23.206 110.065 1.00 18.45
MOTA	1800 N ASP A 228	59.880 22.862 111.018 1.00 21.29
ATOM	1801 CA ASP A 228	59.357 23.032 112.375 1.00 25.80
MOTA	1802 CB ASP A 228	60.464 22.893 113.426 1.00 24.02
ATOM	1803 CG ASP A 228	61.110 21.520 113.422 1.00 26.48
MOTA	1804 OD1 ASP A 228	60.410 20.530 113.135 1.00 29.55
MOTA	1805 OD2 ASP A 228	62.311 21.425 113.744 1.00 29.88
ATOM	1806 C ASP A 228	58.628 24.341 112.620 1.00 27.83
MOTA	1807 O ASP A 228	57.589 24.360 113.284 1.00 25.68
ATOM	1808 N ASN A 229	59.167 25.437 112.098 1.00 25.78
ATOM	1809 CA ASN A 229	58.537 26.739 112.297 1.00 27.75
ATOM	1810 CB ASN A 229	59.453 27.850 111.770 1.00 32.77
ATOM	1811 CG ASN A 229	60.707 28.020 112.621 1.00 30.35
ATOM	1812 OD1 ASN A 229	60.635 28.433 113.782 1.00 33.12
MOTA MOTA	1813 ND2 ASN A 229 1814 C ASN A 229	61.856 27.691 112.053 1.00 28.11
ATOM		57.168 26.817 111.645 1.00 29.47
ATOM		56.230 27.387 112.202 1.00 26.75
ATOM .		57.041 26:228 110.463 1.00 30.80
ATOM	1818 CB GLU A 230	55.761 26.244 109.773 1.00 30.77
ATOM	1819 CG GLU A 230	55.929 25.716 108.341 1.00 29.11 56.897 26.531 107.507 1.00 35.94
ATOM	1820 CD GLU A 230	
ATOM	1821 OE1 GLU A 230	
MOTA	1822 OE2 GLU A 230	
ATOM	1823 C GLU A 230	
MOTA	1824 O GLU A 230	54.723 25.407 110.527 1.00 30.13 53.563 25.799 110.631 1.00 28.35
ATOM	1825 N PHE A 231	55.141 24.262 111.060 1.00 32.49
ATOM	1826 CA PHE A 231	54.223 23.386 111.790 1.00 28.54
ATOM	1827 CB PHE A 231	54.913 22.075 112.191 1.00 31.22
MOTA	1828 CG PHE A 231	53.974 21.050 112.781 1.00 28.41
MOTA	1829 CD1 PHE A 231	53.026 20.417 111.982 1.00 29.66
ATOM	1830 CD2 PHE A 231	54.036 20.723 114.130 1.00 28.38
MOTA	1831 CE1 PHE A 231	52.153 19.469 112.518 1.00 25.79
ATOM	1832 CE2 PHE A 231	53.166 19.774 114.681 1.00 31.40
ATOM	1833 CZ PHE A 231	52.223 19.146 113.870 1.00 30.51
ATOM	1834 C PHE A 231	53.693 24.065 113.045 1.00 26.85
ATOM	1835 O PHE A 231	52.483 24.092 113.277 1.00 25.99
MOTA	1836 N LEU A 232	54.598 24.607 113.858 1.00 26.75
MOTA	1837 CA LEU A 232	54.193 25.283 115.092 1.00 27.25
ATOM	1838 CB LEU A 232	55.422 25.617 115.933 1.00 25.15
ATOM	1839 CG LEU A 232	56.176 24.372 116.420 1.00 28.11
ATOM	1840 CD1 LEU A 232	57.440 24.783 117.162 1.00 27.32
ATOM	1841 CD2 LEU A 232	55.268 23.540 117.328 1.00 27.87
ATOM	1842 C LEU A 232	53.371 26.542 114.800 1.00 26.98
ATOM	1843 O LEU A 232	52.449 26.866 115.544 1.00 23.34
ATOM	1844 N PHE A 233	53.694 27.232 113.708 1.00 24.99
MOTA	1845 CA PHE A 233	52.950 28.426 113.312 1.00 28.13
ATOM	1846 CB PHE A 233	53.542 29.029 112.029 1.00 30.77
ATOM	1847 CG PHE A 233	52.719 30.151 111.448 1.00 29.65
ATOM	1848 CD1 PHE A 233	52.803 31.441 111.962 1.00 32.80

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ATOM	1849	CD	2. PHE A :	233	51.825	29.903	110.412	7.00	31.59
ATOM	1850				52.008		111.452		33.90
MOTA	1851	CE.	2 PHE A	233	51.022	30.924	109.895	1.00	32.47
			PHE A		51.114				32.50
ATOM	1852						110.415		
ATOM	1853	С	PHE A	233	51.510	27.999	113.031	1.00	31.62
	1854		PHE A	222	50.553	28.603	113.532	1 00	25.88
MOTA									
ATOM	1855	N	ALA A	234	51.370	26.955	112.215	1.00	28.12
	1856		ALA A	234	50.056	26 436	111.853	. 1 00	25.68
MOTA									
ATOM	1857	CB	ALA A	234	50.195	25.279	110.864	1.00	20.08
	1858		ALA A	234	49.304	25.969	113.089	1 00	25.17
MOTA									
MOTA	1859	0	ALA A	234	48.114	26.228	113.234	7.00	25.21
MOTA	1860	N	LEU A 2	235	50.002	25.285	113.987	1.00	28.18
MOTA	1861	CA	LEU A 2	435	49.367		115.195		33.70
MOTA	1862	CB	LEU A 2	235	50.356	23.964	116.026	1.00	32.70
	1863	CG	LEU A 2		49.772	22 700	116.820	1 00	36.89
MOTA									
MOTA	1864	CD:	l LEU A 2	235	50.634	22.545	118.052	1.00	31.37
MOTA	1865	CD2	LEU A 2	775	48.344	23 072	117.231	1.00	31.47
ATOM	1866	С	LEU A 2	:35	48.841		116.062	1.00	33.38
ATOM	1867	0	LEU A 2	235	47.673	25.926	116.455	1.00	28.13
MOTA	1868	N	GLU A 2		49.710	26.888			34.02
ATOM	1869	CA	GLU A 2	236	49.336	28.026	117.199	1.00	37.30
	1870	CB	GLU A 2	3.6	50.528	28 972	117.400	1 00	41.51
MOTA									
MOTA	1871	ÇG	GLU A 2	236	51.675	28.356	118.188	1.00	49.54
ATOM	1872	CD	GLU A 2	36	52.811	29.334	118.451	1.00	55.02
ATOM	1873	OE:			53.781	28.947	119.140		56.19
MOTA	1874	OE2	2 GLU A 2	36	52.735	30.486	117.968	1.00	54.84
	1875	C	GLU A 2		48.163	28.803		1 00	33.98
MOTA									
MOTA	1876	0	GLU A 2	36 ⋅	47.211	29.098	117.362	1.00	37.01
	1877	N	LYS A 2	37	48.223	29 137	115.354	1 00	33.94
ATOM									
ATOM	1878	CA	LYS A 2	3/	47.140	29.888	114.726		33.10
ATOM	1879	CB	LYS A 2	37	47.505	30.244	113.281	1.00	36.08
					48.695		113.165		33.62
MOTA	1880	CG	LYS A 2						
ATOM	1881	CD	LYS A 2	37	48.395	32.508	113.856	1.00	37.99
	1882	CE	LYS A 2	37	49.569	33 - 471	113.762	1.00	45.24
ATOM									
ATOM	1883	NZ	LYS A 2	37	49.285		114.500		43.49
ATOM	1884	С	LYS A 2	37	45.820	29.128	114.751	1.00	31.40
			LYS A 2		44.793		115.131		31.67
MOTA	1885	0							
MOTA	1886	N	SER A 2	38	45.841	27.861	114.354		28.72
	1887	CA	SER A 2	3.8	44.610	27 080	114.335	1.00	31.74
ATOM									
ATOM .	1888	CB	SER A 2	38	44.834		113.660		28.90
ATOM	1889	OG	SER A 2	38	45.760	24.924	114.372	1.00	25.18
		c	SER A 2		44.041	26.891			33.23
ATOM	1890								
MOTA	1891	0	SER A 2	38	42.823	26.875	115.916	1.00	34.79
ATOM	1892	N	LEU A 2	39	44.907	26.742	116.741	1.00	35.27
									37.57
MOTA	1893	CA	LEU A 2		44.413	26.587			
MOTA	1894	CB	LEU A 2	39	45.554	26.307	119.090	1.00	38.58
	1895	CG	LEU A 2	30 .	46.176	24.907	119.038	1.00	39.74
MOTA									
ATOM	1896	CD1	LEU A 2	39	47.276	24.797	120.075		35.82
ATOM	1897	CD2	LEU A 2	39	45.109	23.861	119.301	1.00	34.93
_							118.521	1 00	39.09
MOTA	1898	С	LEU A 2		43.670	27.852			
ATOM	1899	0	LEU A 2	39	42.628	27.782	119.174	1.00	35.50
	1900	N.T	GLU A 2	40	44.202	29.007	118.131	1 00	39.27
ATOM		N	GEO A 2	40					
ATOM	1901	CA	GLU A 2	40	43.561	30.281	118.450	1.00	40.15
	1902	CB	GLU A 2	40	44.366	31.448	117.883	1.00	40.42
MOTA									
MOTA	1903	CG	GLU A 2		45.661	31.747			43.91
MOTA	1904	CD	GLU A 2	40	46.407	32.884	117.942	1.00	49.31
			GLU A 2		45.772	33.925	117.665		49.00
MOTA	1905								
ATOM	1906	OE2	GLU A 2	40	47.624	32.745			54.05
	1907	C	GLU A 2		42.165	30.312	117.849	1.00	39.58
ATOM									
MOTA	1908	0	GLU A 2		41.224		118.455	1.00	
ATCM	1909	N	ILE A 2	41	42.039		116.645	1.00	35.70
			ILE A 2		40.754		115.964	1.00	
MOTA	1910	CA							
ATOM	1911	CB	ILE A 2	41	40.904		114.546	1.00	
	1912	CG2	ILE A 24	41	39.535	29.005	113.895	1.00	37.30
ATOM			TIE NO	17			113.724	1.00	
ATOM	1913	CG1	ILE A 24	± 4.	41.832	30.048			
ATOM	1914	CD1	ILE A 24	11	42.106	29.541	112.320	1.00	36.1 5

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ATOM	1915 C ILE A 241	39.75	1 28.881 116.737	1.00 37.31
ATOM	1916 O ILE A 241	38.59		
ATOM	1917 N VAL A 242	40.20		
ATOM	1918 CA VAL A 242	39.33		
MOTA	1919 CB VAL A 242	40.025		
MOTA	1920 CG1 VAL A 242	39.120		
MOTA	1921 CG2 VAL A 242	40.364		
MOTA	1922 C VAL A 242	38.930	27.451 119.305	1.00 40.40
MOTA	1923 O VAL A 242	37.759	27.422 119.675	1.00 38.19
ATOM	1924 N LYS A 243	39.905		1.00 45.47
ATOM	1925 CA LYS A 243 .	39.661	-	1.00 48.74
ATOM	1926 CB LYS A 243	40.945		1.00 51.01
MOTA	1927 CG LYS A 243	41.853		1.00 56.09
MOTA	1928 CD LYS A 243 1929 CE LYS A 243	41.250		1.00 57.39
MOTA	1929 CE LYS A 243 1930 NZ LYS A 243	41.054		1.00 59.53
MOTA MOTA	1930 NZ LIS A 243	40.448		1.00 57.91
ATOM	1932 O LYS A 243	38.559		1.00 51.67
MOTA	1933 N GLU A 244	37.815 38.451		1.00 52.84
MOTA	1934 CA GLU A 244	37.460		1.00 53.77
ATOM	1935 CB GLU A 244	37.954		1.00 54.74 1.00 55.15
ATOM	1936 CG GLU A 244	37.068	33.717 118.865	1.00 60.63
ATOM	1937 CD GLU A 244	37.602	34.714 117.868	1.00 65.87
ATOM	1938 OE1 GLU A 244	38.746	35.181 118.053	1.00 70.36
MOTA	1939 OE2 GLU A 244	36.879	35.031 116.900	1.00 67.09
MOTA	1940 C GLU A 244	36.051	31.025 119.626	1.00 52.65
MOTA	1941 O GLU A 244	35.127	31.838 119.606	1.00 55.59
MOTA	1942 N VAL A 245	35.869	29.745 119.332	1.00 50.57
ATOM	1943 CA VAL A 245	34.546	29.269 118.947	1.00 45.75
ATOM	1944 CB VAL A 245	34.475	29.081 117.409	1.00 46.91
MOTA	1945 CG1 VAL A 245	33.085	28.634 116.986	1.00 52.62
MOTA	1946 CG2 VAL A 245	34.825	30.389 116.716	1.00 48.34
MOTA	1947 C VAL A 245 1948 O VAL A 245	34.130	27.969 119.642	1.00 42.67
MOTA MOTA	1948 O VAL A 245 1949 N PHE A 246	33.021	27.480 119.445	1.00 43.61
ATOM	1950 CA PHE A 246	35.001 34.662	27.417 120.477 26.168 121.139	1.00 40.87
MOTA	1951 CB PHE A 246	35.106	24.991 120.257	1.00 37.47
ATOM	1952 CG PHE A 246	34.450	23.685 120.604	1.00 37.00 1.00 33.22
ATOM	1953 CD1 PHE A 246	33.111	23.467 120.302	1.00 33.22
ATOM	1954 CD2 PHE A 246	35.168	22.674 121.234	1.00 32.13
MOTA	1955 CE1 PHE A 246	32.493	22.260 120.621	1.00 37.75
MOTA	1956 CE2 PHE A 246	34.561	21.459 121.561	1.00 35.92
MOTA	1957 CZ PHE A 246	33.217	21.252 121.251	1.00 36.30
ATOM	1958 C PHE A 246	35.322	26.065 122.509	1.00 38.93
MOTA	1959 O PHE A 246	36.546	26.158 122.630	1.00 40.66
MOTA	1960 N GLU A 247	34.500	25.870 123.537	1.00 38.59
MOTA	1961 CA GLU A 247	34.970	. 5.733 124.918	1.00 44.60
ATOM	1962 CB GLU A 247 1963 CG GLU A 247	34.146	5.615 125.865	1.00 47.07
MOTA	1963 CG GLU A 247 1964 CD GLU A 247	33.161	27.569 125.185	1.00 56.16
ATOM ATOM	1965 OE1 GLU A 247	31.944	26.865 124.577 26.088 123.607	1.00 62.03
ATOM	1966 OE2 GLU A 247	30.822	27.094 125.083	
ATOM	1967 C GLU A 247	34.774	24.269 125.285	1.00 64.59
ATOM	1968 O GLU A 247	33.727	23.879 125.794	1.00 39.91
MOTA	1969 N PRO A 248	35.792	23.442 125.041	1.00 38.64
ATOM	1970 CD PRO A 248	37.101	23.817 124.483	1.00 33.25
	1971 CA PRO A 248	35.769	22.006 125.316	1.00 35.84
MOTA	1972 CB PRO A 248	37.047	21.531 124.648	1.00 36.05
MOTA	1973 CG PRO A 248	37.970	22.687 124.982	1.00 34.21
ATOM	1974 C PRO A 248	35.736	21.611 126.779	1.00 33.94
ATOM	1975 0 PRO A 248	36.445	22.186 127.597	1.00 32.05
atom	1976 N GLU A 249	34.914	20.616 127.096	1.00 29.39
ATOM	1977 CA GLU A 249	34.841	20.105 128.459	1.00 33.12
MOTA	1978 CB GLU A 249	33.521	19.361 128.693	1.00 30.36
ATCM	1979 CG GLU A 249	32.284	20.212 128.564	1.00 35.98
atcm	1980 CD GLU A 249	31.026	19.388 128.668	1.00 40.52

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```
MOTA
        1981
              OE1 GLU A 249
                                   30.817
                                           18.509 127.804
                                                             1.00 40.27
        1982
              OE2
                  GLU A 249
                                   30.252
                                           19.609 129.620
                                                             1.00 38.57
ATOM
                                                             1.00 32.30
ATOM
        1983
              C
                   GLU A 249
                                   35.995
                                            19.119 128.623
        1984
                   GLU A 249
                                   36.472
                                            18.884 129.728
                                                             1.00 28.51
ATOM
        1985
                   VAL A 250
                                   36.434
                                            18.547 127.502
                                                             1.00 33.74
              N
MOTA
                                                             1.00 29.31
                   VAL A 250
                                   37.516
                                            17.563 127.494
ATOM.
        1986
              CA
        1987
              CB
                   VAL A 250
                                   36.988
                                            16.174 127.926
                                                             1.00 29.85
ATOM
                   VAL A 250
                                   35.908
                                           15.711 126.958
                                                             1.00 24.36
        1988
              CG1
ATOM
                                            15.166 127.978
                                                             1.00 25.60
MOTA
        1989
              CG2
                  VAL A 250
                                   38.121
        1990
                  _VAL A 250
                                   38.066
                                           17.453 126.076
                                                             1.00 29.30
              С
ATOM
                                           17.741 125.114
                                                             1.00 24.46
                   VAL A 250
                                   37.358
ATOM
        1991
              0
                                                             1.00 27.96
MOTA
        1992
              N
                   TYR A 251
                                   39.323
                                           17.046 125.930
                   TYR A 251
                                   39.865
                                           16.913 124.585
                                                             1.00 30.06
        1993
              CA
ATOM
                                           18.206 124.165
                                   40.585
                                                             1.00 25.89
                   TYR A 251
        1994
              CB
ATOM
                                                             1.00 29.90
MOTA
        1995
              CG
                   TYR A 251
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                                           18.370 124.692
                  TYR A 251
                                   43.087
                                           17.794 124.029
                                                             1.00 26.02
        1996
              CD1
ATOM
                  TYR A 251
                                   44.390
                                           17.953 124.507
                                                             1.00 29.20
              CE1
        1997
ATOM
ATOM
        1998
              CD2
                  TYR A 251
                                   42.249
                                           19.107 125.849
                                                             1.00 31.96
                  TYR A 251
                                   43.551
                                           19.271 126.338
                                                             1.00 31.54
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              CE2
MOTA
                   TYR A 251
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                                           18.694 125.664
                                                             1.00 31.46
        2000
              CZ.
ATOM
                                                             1.00 29.69
                                           18.854 126.152
MOTA
        2001
              OH
                  TYR A 251
                                   45.894
                   TYR A 251
                                   40.801
                                           15.731 124.451
                                                             1.00 27.56
MOTA
        2002
                                   41.382
                                                             1.00 28.23
       2003
              0
                  TYR A 251
                                           15.273 125.436
MOTA
                                   40.908
                                           15.222 123.227
                                                             1.00 23.52
ATOM
       2004
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                  LEU A 252
        2005
              CA
                  LEU A 252
                                   41.806
                                           14.117 122.919
                                                             1.00 26.53
MOTA
                                   41.057
                                           12.930 122.293
                                                             1.00 25.74
       2006
              CB
                  LEU A 252
ATOM
                                   40.266
                                           12.001 123.221
                                                             1.00 28.49
                  LEU A 252
ATOM
       2007
              CG
                                           12.753 123.868
                                                             1.00 27.67
       2008
              CD1
                  LEU A 252
                                   39.122
ATOM
                  LEU A 252
                                   39.727
                                           10.835
                                                  122.414
                                                             1.00 32.00
       2009
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MOTA
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                  LEU A 252
                                   42.842
                                           14.638 121.932
MOTA
       2010
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                                           15.444 121.055
                                                             1.00 24.42
                  LEU A 252
                                   42.528
ATOM
       2011
              O
                  LEU A 253
                                   44.075
                                           14.176
                                                  122.078
                                                             1.00 24.60
ATOM
       2012
              N
                                   45.157
                                           14.599 121.204
                                                             1.00 25.04
       2013
                  LEU A 253
MOTA
              CA
                                           15.400 122.017
                                                             1.00 22.48
                  LEU A 253.
                                   46.176
ATOM
       2014
              CB
                                                             1.00 21.05
                                           15.880 121.323
ATOM
       2015
              CG
                  LEU A 253
                                   47.456
                  LEU A 253
                                                             1.00 23.05
                                   47.105
                                           16.833 120.175
       2016
              CD1
ATOM
                                                             1.00 16.40
       2017
              CD2
                  LEU A 253
                                   48.348
                                           16.578 122.360
ATOM
                                   45.822
                                           13.374 120.580
                                                             1.00 23.55
                        253
ATOM
       2018
              C
                  LEU A
                                                             1.00 22.11
       2019
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                                   46.329
                                           12.516 121.303
ATOM
                  GLN A 254
                                   45.811
                                           13.287 119.248
                                                             1.00 22.33
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             N
MOTA
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                                                             1.00 19.84
                  GLN A 254
                                   46.417
       2021
             CA
MOTA
                                                             1.00 23.09
                  GLN A 254
                                           11.731 117.348
                                   45.542
ATOM
       2022
             CB
                                                             1.00 35.49
                  GLN A 254
                                   46.075
                                           12.038 115.963
       2023
             CG
ATOM
             CD
                  GLN A 254
                                   47.073
                                           11.017 115.453
                                                             1.00 31.26
       2024
ATOM
                                            9.937 114.961
                                                             1.00 33.69
                                   46.712
                  GLN A 254
       2025
             OE1
ATOM
                                                             1.00 31.02
             NE2
                  GLN A 254
                                   48.338
                                           11.349 115.574
       2026
: TOM
                  GLN A 254
                                   47.831
                                           12.576 118.153
                                                             1.00 22.46
       2027
             С
..TOM
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                                           13.599 117.478
                                                             1.00 17.56
                  GLN A 254
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                                           11.781 118.590
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                  LEU A 255
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ATOM
                  LEU
                      A 255
                                   50.213
                                           12.079 118.383
                                                             1.00 17.04
       2030
             CA
MOTA
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                                           12.136 119.750
                                                             1.00 14.75
       2031
             CB
ATOM
                                                             1.00 25.02
                  LEU A 255
                                  50.277
                                           13.196 120.670
ATOM
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             CD1
                 LEU A 255
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                                           12.996 122.107
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MOTA
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                                                             1.00 18.30
                 LEU A 255
                                           14.578 120.149
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       2034
             CD2
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                                           11.169 117.476
                  LEU A 255
       2035
             С
                                  51.023
MOTA
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                                           10.705 117.875
                                                             1.00 18.73
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MOTA
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                                           10.928 116.259
                                                             1.00 22.75
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ATOM
       2037
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                                                             1.00 24.09
                                           10.093 115.330
                 GLY A 256
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                                                             1.00 24.27
                  GLY A 256
                                  52.660
                                           10.721 115.126
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             C
ATOM
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                                           11.945 115.134
                                                             1.00 19.15
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MOTA
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                                  53.680
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                  THR A 257
ATOM
                                                             1.00 21.32
                  THR A 257
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                                           10.440 114.765
             CA
MOTA
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                                                             1.00 17.24
                 THR A 257
                                  56.048
       2043
             CB
MOTA
                                            8.248 115.004
                                                            1.00 17.48
                 THR A 257
       2044
             0G1
                                  56.009
ATOM
                                                            1.00 15.73
                                            9.532 117.016
                 THR A 257
                                  55.728
       2045
             CG2
MOTA
                  THR A 257
                                  55.403
                                           10.527 113.290
                                                            1.00 22.98
       2046
ATOM
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```
MOTA
          2047
                0
                     THR A 257
                                     56.517
                                             10.941 112.974
                                                               1.00 20.39
  MOTA
          2048
                N
                     ASP A 258
                                     54.493
                                              10.177 112.379
                                                               1.00 22.20
 ATOM
          2049
                CA
                     ASP A 258
                                     54.863
                                             10.229 110.961
                                                               1.00 26.56
 ATOM
          2050
                CB
                     ASP A 258
                                     53.849
                                               9.496
                                                     110.056
                                                               1.00 25.06
          2051
 ATOM
                CG
                     ASP A 258
                                     52.415
                                               9.944 110.252
                                                               1.00 29.08
 ATOM
          2052
                С
                     ASP A 258
                                     55.222
                                             11.596 110.364
                                                               1.00 27.87
                     ASP A 258
 MOTA
          2053
                O
                                     55.756
                                             11.661 109.254
                                                               1.00 25.61
         2054
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                                                               1.00 29.86
 MOTA
          2055
                OD2
                    ASP A 258
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                                                               1.00 33.25
         2056
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                    PRO A 259
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                    PRO A 259
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                                                               1.00 28.59
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                CA
                    PRO A 259
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                                             14.006 110.469
                                                               1.00 30.95
 MOTA
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                CB
                    PRO A 259
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                                             14.993
                                                     111.296
                                                               1.00 35.06
         2060
                CG
 MOTA
                    PRO A 259
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                                             14.320
                                                     112.636
                                                               1.00 35.26
                                    56.790
 MOTA
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                                                               1.00 29.06
                    PRO A 259
 MOTA
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                                                    111.280
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                                             13.545 111.455
                                                               1.00 28.41
                    LEU A 260
 ATOM
         2065
                CB
                                    59.461
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                                                               1.00 22.47
         2066
 MOTA
               CG
                    LEU A 260
                                    58.970
                                             12.791 113.969
                                                               1.00 20.14
 ATOM
         2067
               CD1
                    LEU A 260
                                    59.352
                                             11.599 114.826
                                                               1.00 22.83
 MOTA
         2068
               CD2
                   LEU A 260
                                    59.592
                                             14.079
                                                    114.532
                                                              1.00 20.48
         2069
 ATOM
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                    LEU A 260
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                                                              1.00 27.95
 ATOM
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                    LEU A 260
                                    59.407
                                             12.535 109.299
                                                              1.00 27.03
         2071
 ATOM
               N
                    LEU A 261
                                    60.874
                                             14.081 110.040
                                                              1.00 26.76
         2072
               CA
 ATOM
                   LEU A 261
                                    61.742
                                             14.010 108.865
                                                              1.00 26.56
 MOTA
         2073
               CB
                   LEU A 261
                                    63.067
                                             14.737 109.137
                                                              1.00 23.06
                                    64.131
63.642
         2074
 MOTA
               CG
                   LEU A 261
                                             14.615
                                                    108.025
                                                              1.00 29.52
         2075
 MOTA
               CD1
                   LEU A 261
                                            15.325 106.770
                                                              1.00 22.68
        2076
 ATOM
               CD2
                   LEU A 261
                                    65.460
                                            15.219
                                                    108.475
                                                              1.00 26.71
        2077
                   LEU A 261
                                            12.577
 ATOM
               C
                                    62.063
                                                    108.443
                                                              1.00 28.23
 MOTA
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               0
                   LEU A 261
                                    61.880
                                            12.198 107.289
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ATOM	2115	O LEU A 265	53.457	9.796 104.979	1.00 21.30
ATOM	2116	N SER A 266	55.540	10.622 104.959	1.00 23.23
MOTA	2117		55.084		1.00 26.30
ATOM	2118		54.856	12.502 106.444	1.00 25.16
ATOM	2119	OG SER A 266	56.074	12.845 107.084	1.00 22.92
ATOM	2120	C SER A 266	56.147		1.00 30.17
	2121	O SER A 266	57.334		
MOTA					1.00 31.65
MOTA	2122	N LYS A 267	55.731	13.985 103.757	1.00 31.56
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ATOM	2124	CB LYS A 267	56.140	15.425 101.834	1.00 30.54
	2125	CG LYS A 267			
ATOM			55.815	14.327 100.819	1.00 34.13
ATOM	2126	CD LYS A 267	57.039	13.463 100.549	1.00 29.09
ATOM	2127	CE LYS A 267	56.745	12.376 99.524	1.00 37.61
MOTA	2128	NZ LYS A 267	57.956	11.541 99.272	1.00 31.91
	2129	C LYS A 267			
ATOM			57.050	16.004 104.107	1.00 30.85
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MOTA	2131	N PHE A 268	56.688	15.826 105.377	1.00 24.19
ATOM	2132	CA PHE A 268	57.009	16.808 106.412	1.00 25.34
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ATOM	2134	CG PHE A 268	54.636	17.256 107.257	1.00 21.68
ATOM	2135	CD1 PHE A 268	53.631	17.221 108.216	1.00 28.65
ATOM	2136	CD2 PHE A 268	54.346	17.806 106.011	1.00 25.14
	2137	CE1 PHE A 268			
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MOTA	2138	CE2 PHE A 268	53.077	18.315 105.730	1.00 23.79
MOTA	2139	CZ PHE A 268	52.082	18.275 106.702	1.00 28.13
ATOM	2140	C PHE A 268	58.410	16.470 106.908	1.00 25.66
	2141	O PHE A 2.68	58.778		
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MOTA	2142	n asna 269	59.194	17.490 107.230	1.00 25.81
MOTA	2143	CA ASN A 269	60.555	17.270 107.709	1.00 30.60
ATOM	2144	CB ASN A 269	61.566	17.938 106.767	1.00 31.97
	2145	CG ASN A 269 ·	61.392	17.513 105.317	1.00 35.83
ATOM					
MOTA	2146	OD1 ASN A 269	61.235	16.332 105.020	1.00 33.93
ATOM	2147	ND2 ASN A 269	61.446	18.477 104.405	1.00 33.95
ATOM	2148	C ASN A 269	60.723	17.843 109.110	1.00 31.80
ATOM	2149	O ASN A 269	61.609	18.665 109.348	1.00 28.13
ATOM	2150	N LEU A 270	59.888	17.397 110.043	1.00 29.70
MOTA	2151	CA LEU A 270	59.954	17.918 111.406	1.00 26.87
ATOM	2152	CB LEU A 270	58.575	17.833 112.074	1.00 26.60
ATOM	2153	CG LEU A 270	57.392	18.425 111.297	1.00 29.62
	2154	CD1 LEU A 270	56.177		
MOTA				18.494 112.222	1.00 28.54
ATOM	2155	CD2 LEU A 270	57.740	19.825 110.790	1.00 29.40
ATOM	2156	C LEU A 270	60.979	17.242 112.301	1.00 26.83
ATOM	2157	O LEU A 270	61.490	16.158 111.990	1.00 19.60
ATOM	2158	N SER A 271	61.275	17.896 113.420	1.00 21.66
ATOM	2159	CA SER A 271	62.220	17.365 114.393	1.00 27.08
MOTA	2160	CB SER A 271	63.189	18.460 114.846	1.00 24.64
ATOM	2161	OG SER A 271	62.499	19.433 115.626	1.00 18.60
ATOM	2162	C SER A 271	61.454	16.868 115.618	1.00 23.70
MOTA	2163	O SER A 271	60.272	17.150 115.772	1.00 22.56
MOTA	2164	n asna 272	62.157	16.129 116.470	1.00 28.35
ATOM	2165	CA ASN A 272	61.649	15.593 117.739	1.00 31.03
ATOM	2166	CB ASN A 272	62.774	14.880 118.498	1.00 28.80
MOTA	2167		62.854	13.428 118.180	1.00 36.65
MOTA	2168	OD1 ASN A 272	63.712	12.709 118.705	1.00 29.89
MOTA	2169	ND2 ASN A 272	61.953	12.969 117.319	1.00 40.80
ATOM	2170	C ASN A 272	61.167	16.695 118.661	1.00 31.83
ATOM	2171		60.090	16.618 119.261	1.00 27.50
ATOM	2172	N VAL A 273	62.032	17.693 118.804	1.00 31.49
ATOM	2173	CA VAL A 273	61.802	18.837 119.667	1.00 31.58
ATOM	2174	CB VAL A 273	63.069	19.709 119.725	1.00 35.80
	2175	CG1 VAL A 273	62.804		
ATOM				20.988 120.500	1.00 48.07
ATOM	2176	CG2 VAL A 273	64.198	18.914 120.381	1.00 42.81
ATCM	2177	C VAL A 273	60.608	19.665 119.234	1.00 30.13
ATOM	2178	O VAL A 273	59.872	20.174 120.072	1.00 31.44

1OTA	1 2179 N ALA A 274	60.40	5 19.800 117.92	
ATO1		59.25		
ATON				
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ATOM		58.00		
		56.96		
ATOM		58.122		0 1.00 25.20
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ATOM		- 56.365		
ATOM		54.248		
· ATOM		24.240		
ATOM	2192 CZ PHE A 275	55.343	13.231 119.05	
ATOM		54.282		
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ATOM		55.428		7 1.00 22.40
MOTA		57.594		1 1.00 25.45
MOTA		57.357	17.837 121.766	5 1.00 27.94
ATOM		58.667	17.692 122.534	1.00 26.11
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MOTA	2206 CG LYS A 277	. 57.941	23.934 121.237	1.00 36.46
ATOM	2207 CD LYS A 277	56.633	24.668 121.454	
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ATOM	2213 CA ALA A 278	53.662	21.096 119.676	1.00 30.15
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ATOM	2215 C ALA A 278	52.789	20.270 118.406	
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ATOM	2219 CB PHE A 279	53.343	17.534 122.886	1.00 26.83
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MOTA	2223 CE1 PHE A 279	51.054	15.500 125.121	1.00 37.90
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ATOM	2225 CZ PHE A 279	51.783	15.469 126.313	1.00 36.23
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ATOM	2243 O ILE A 281		24.621 123.854	1.00 30.15
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MOTA
                                                             1.00 30.98
                  GLU A 288
                                   44.309
                                            17.623 129.546
MOTA
        2300
              С
                                                             1.00 28.67
                  GLU A 288
                                   43.144
                                            17.957 129.762
ATOM
        2301
              0
                                   44.641
                                            16.694 128.657
                                                             1.00 29.66
MOTA
        2302
                  GLY A 289
              N
                                                             1.00 28.08
                  GLY A 289
                                   43.625
                                            15.999 127.886
        2303
              CA
MOTA
                  GLY A 289
                                   43.922
                                            14.510 127.861
                                                             1.00 30.45
ATCM
        2304
              C
                                   44.618
                                            14.307 128.739
                                                             1.00 25.90
        2305
                  GLY A 289
ATOM
                  VAL A 290
                                   43.384
                                            13.807 126.868
                                                             1.00 26.21
        2306
              N
ATOM
                                                             1.00 27.31
                  VAL A 290
                                   43.612
                                            12.373 126.718
        2307
              CA
ATOM
                                   42.288
                                                             1.00 26.53
                                            11.626 125.412
                  VAL A 290
        2308
              CB
ATOM
       2309
              CG1 VAL A 290
                                   42.554
                                            10.148 126.204
                                                             1.00 25.36
ATOM
                                                             1.00 24.38
                  VAL A 290
                                   41.308
                                           11.822 127.565
        2310
              CG2
ATOM
```

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ATOM	2311	1 (VAL A 290	44.580	12.2	48 125.55	0 1.00 24.52
ATOM	2312	2 C	VAL A 290	44.307	12.7	43 124.46	1 1.00 26.84
ATOM	2313	3 N	TYR A 291	45.716	11 5	97 125.77	5 1.00 23.56
ATOM	2314	1 C	A TYR A 291	46.729	11 4	78 124.73	2 1.00 23.74
ATOM	2315	5 C	B TYR A 291	48.092	11 8	17 125.34	
ATOM	2316		G TYR A 291	48.040			
ATOM	2317		D1 TYR A 291	48.326			
ATOM			E1 TYR A 291				
ATOM			D2 TYR A 291	48.200		1 128.214	
MOTA	2320		E2 TYR A 291	47.634			
MOTA	2321			47.504	15.47		
ATOM	2322			47.786			
ATOM	2323			47.631	16.65	7 128.283	
	2323		TYR A 291	46.768	10.11	.8 124.044	
MOTA	2325		TYR A 291	46.837		124.707	
ATOM				46.755	10.14		
ATOM	2326			46.767	8.92	4 121.902	
ATOM	2327	CI		45.482	8.84	2 121.076	1.00 22.13
MOTA	2328	CC		44.162	9.06		1.00 23.78
ATOM	2329		01 LEU A 292	43.001	8.95	9 120.826	1.00 23.09
ATOM	2330	CI		44.008	8.05	0 122.930	1.00 16.01
ATOM	2331	С	LEU A 292	47.953	8.88		
MOTA	2332	0	LEU A 292	48.527	9.92		
ATOM	2333	N	GLY A 293	48.301	7.68	4 120.491	1.00 18.83
ATOM	2334	CA	GLY A 293	49.401	7.52	9 119.554	1.00 24.35
ATOM	2335	C	GLY A 293	48.908		6 118.154	1.00 24.22
MOTA	2336	0	GLY A 293	48.025		4 117.991	1.00 20.46
ATOM	2337	N	GLY A 294	49.459	7.17		1.00 24.63
ATOM	2338	CA	GLY A 294	49.035	7.42		1.00 22.03
ATOM	2339	С	GLY A 294	50.024		9 114.769	1.00 22.90
ATOM	2340	0	GLY A 294	50.956	6.15	0 115.136	1.00 24.10
ATOM	2341	N	GLY A 295	49.825	7.20		1.00 19.85
MOTA	2342	CA	GLY A 295	50.721	6.72		1.00 23.33
ATOM	2343	С	GLY A 295	52.185		112.740	1.00 19.01
ATOM	2344	0	GLY A 295 .	52.541	8.094		1.00 19.39
MOTA	2345	N	GLY A 296	53.035		112.472	1.00 25.85
ATOM	2346	CA	GLY A 296	54.468		2 112.690	1.00 22.65
MOTA	2347	С	GLY A 296	55.098	4.898	3 112.146	1.00 25.31
ATOM	2348	0	GLY A 296	54.778	3.798	3 112.609	1.00 25.86
ATOM	2349	N	TYR A 297	56.005	5.034	111.185	1.00 22.83
ATOM	2350	ÇA	TYR A 297	56.598	3.852		1.00 23.93
ATOM	2351	CB	TYR A 297	56.137	3.780	109.125	1.00 21.59
ATOM	2352	CG	TYR A 297	54.660	4.084	109.035	1.00 25.85
ATOM	2353	CD1		54.203	5.402		1.00 22.28
ATOM	2354	CE1		52.842	5.695	109.089	1.00 20.19 [.]
ATOM	2355	CD2	 .	53.713	3.062		1.00 25.28
ATOM	2356	CE2		52.352	3.346		1.00 21.83
ATOM	2357	CZ	TYR A 297	51.927	_		1.00 21.81
ATOM	2358	OH	TYR A 297	50.588	4.972		1.00 19.51
ATOM	2359	C	TYR A 297	58.104	3.694		1.00 23.78
ATOM	2360	0	TYR A 297	58.665		110.154	1.00 22.07
TOM	2361	N	HIS A 298	58.765		111.326	1.00 23.71
TOM		CA	HIS A 298	60.204	4.534	111.517	1.00 26.33
ATOM		CB	HIS A 298	60.913	5.852	111.216	1.00 28.74
MOTA		CG	HIS A 298	62.403	5.727	111.213	1.00 33.08
MOTA			HIS A 298	63.273	5.465	112.215	1.00 31.83
TOM			HIS A 298	63.151	5.775	110.056	1.00 32.49 ·
TOM			HIS A 298	64.419	5.547	110.345	1.00 29.10
TCM		NE2	HIS A 298	64.520	5.354	111.648	1.00 38.70
TOM		C	HIS A 298	60.371	4.188	112.996	1.00 27.81
TOM		0	HIS A 298	60.120	5.020	113.865	1.00 25.07
TOM.		N	PRO A 299	60.829	2.963	113.297	1.00 29.37
TOM		CD	PRO A 299	61.285		112.352	1.00 26.09
MOT		CA	PRO A 299	61.024		114.669	1.00 29.46
TOM		CB	PRO A 299	61.675		114.465	1.00 29.03
TOM		CG C	PRO A 299 PRO A 299	62.411		113.137	1.00 27.34
TOM	ا ۱۰ دی	-	PRU A 433	61.849	3.403	115.570	1.00 31.88

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Figure 18-37

ATOM	237	7 O PROA 299	61.480	3.622 116.724	1.00 32.45
ATOM	2378		62.959	3.932 115.058	1.00 32.41
MOTA	2379		63.803	4.801 115.878	
ATOM	2380		65.163	5.058 115.207	
ATOM	2381		65.912	3.821 114.738	
ATOM	2382		65.517	2.539 115.120	
ATOM	2383				
MOTA	2384		66.214 67.027	1.411 114.682	1.00 30.68
ATOM	2385			3.941 113.908	
	2386		67.730	2.829 113.466	
ATOM			67.320	1.568 113.854	1.00 33,89
MOTA	2387		68.011	0.471 113.404	
MOTA	2388		63.113	6.134 116.137	1.00 23.44
ATOM	2389		63.108	6.631 117.264	
MOTA	2390 2391		62.530	6.711 115.092	
MOTA			61.839	7.993 115.216	
ATOM	2392		61.266	8.416 113.864	
ATOM	2393		60.715	7.878 116.237	
ATOM	2394		60.556	8.728 117.117	
ATOM	-2395		59.940	6.808 116.110	
ATOM	2396		58.818	6.566 116.996	
ATOM	2397		58.036	5.354 116.483	
ATOM	2398	CG LEU A 302	56.866	4.798 117.291	
MOTA	2399	CD1 LEU A 302	55.983	3.938 116.394	
ATOM	2400	CD2 LEU A 302	57.394	4.001 118.465	
ATOM	2401	C LEU A 302	59.246	6.373 118.451	
ATOM	2402	O LEU A 302	58.648	6.946 119.358	
MOTA	2403	N ALA A 303	60.289	5.580 118.672	1.00 27.85
MOTA	2404 2405	CA ALA A 303	60.765	5.320 120.024	
MOTA	2405	CB ALA A 303	61.854	4.269 119.990	
MOTA	2407	C ALA A 303 O ALA A 303	61.279	6.580 120.714	1.00 26.64
MOTA MOTA	2407		. 60.944	6.849 121.875	1.00 23.18
ATOM	2408		62.092	7.354 120.003	1.00 27.48
ATOM	2410	CA ARG A 304 CB ARG A 304	62.648	8.570 120.581	1.00 25.46
MOTA	2411	CG ARG A 304	63.773 65.005	9.136 119.704	1.00 21.31
MOTA	2412	CD ARG A 304	66.153	8.231 119.562	1.00 25.98
MOTA	2413	NE ARG A 304	65.647	9.042 118.951 9.766 117.796	1.00 27.87
ATOM	2414	CZ ARG A 304	66.207	10.838 117.261	1.00 36.76
ATOM	2415	NH1 ARG A 304	67.323	11.345 117.768	1.00 30.79
ATOM	2416	NH2 ARG A 304	65.623	11.419 116.225	1.00 30.11 1.00 36.07
ATOM	2417	C ARG A 304	61.585	9.634 120.803	1.00 36.07
MOTA	2418	O ARG A 304	61.519	10.237 121.876	1.00 24.23
ATOM	2419	N ALA A 305	60.741	9.854 119.802	1.00 24.23
ATOM	2420	CA ALA A 305	59.700	10.868 119.910	1.00 26.70
ATOM	2421	CB ALA A 305	58.914	10.960 118.607	1.00 28.14
MOTA	2422	C ALA A 305	58.749	10.626 121.072	1.00 25.54
ATOM	2423	O ALA A 305	58 513	11.520 121.883	1.00 24.17
ATOM	2424	N TRP A 306	58 1.89	9.426 121.160	1.00 25.66
ATOM	2425	CA TRP A 306	57.270	9.157 122.253	1.00 28.01
ATOM	2426	CB TRP A 306	56.454	7.873 122.012	1.00 18.66
ATOM	2427	CG TRP A 306	55.382	8.052 120.973	1.00 21.80
ATOM	2428	CD2 TRP A 306	54.709	7.019 120.240	1.00 24.88
ATOM	2429	CE2 TRP A 306	53.725	7.646 119.442	1.00 23.98
ATOM	2430	CE3 TRP A 306	54.839	5.623 120.181	1.00 23.26
MOTA	2431	CD1 TRP A 306	54.795	9.228 120.599	1.00 20.24
ATOM	2432	NE1 TRP A 306	53.799	8.995 119.681	1.00 24.18
MOTA	2433	CZ2 TRP A 306	52.875	6.926 118.590	1.00 24.30
ATOM	2434	CZ3 TRP A 306	53.993	4.906 119.335	1.00 23.89
ATOM	2435	CH2 TRP A 306	53.024	5.562 118.550	1.00 24.12
MOTA	2436	C TKP A 306	57.969	9.113 123.605	1.00 27.93
ATOM	2437	O TRP A 306	57.330	9.319 124.637	1.00 28.58
ATOM	2438	N THR A 307	59.273	8.851 123.615	1.00 26.76
ATOM	2439	CA THR A 307	60.000	8.850 124.881	1.00 22.81
ATOM	2440	CB THR A 307	61.457	8.319 124.730	1.00 25.54
ATOM	2441	OG1 THR A 307	61.435	6.902 124.504	1.00 22.73
ATOM	2442	CG2 THR A 307	62.269	8.599 125.988	1.00 24.03

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ATOM	244:	3 C THR A 307	60.02	7 10.288 12	5.396 1.00 26.	54
ATOM	244	4 O THR A 307	59.92			
ATOM	2445		60.15		4.478 1.00 21.	
ATOM	2446		60.17			
	2447				4.862 1.00 21.	
MOTA			60.44		3.642 1.00 19.	
ATOM	2448		61.79	7 13.386 12	2.938 1.00 21	31
MOTA	2449	9 CD1_LEU A 308	61.90	0 14.362 12	1.774 1.00 21.	75
ATOM	2450	D CD2 LEU A 308	62.93	7 13.622 12	3.915 1.00 19.	
MOTA	2451	L C LEU A 308	58.81	•	5.479 1.00 25.	
ATOM	2452		58.73		6.561 1.00 21.	
	2453					
MOTA			- 57.74		4.806 1.00 21.	
MOTA	2454		56.39		5.298 1.00 19.2	23
MOTA	2455		55.33	7 12.149 12	4.366 1.00 19.0	63
MOTA	2456	CG2 ILE A 309	53.94	5 12.321 12	4.948 1.00 19.	54
ATOM	2457	CG1 ILE A 309	55.40	3 12.788 12	2.979 1.00 20.8	
ATOM	2458		55.11		2.988 1.00 20.0	
ATOM	2459		56.22			
	2460					
MOTA			55.73		7.602 1.00 21.3	
ATOM	2461		56.65		6.888 1.00 26.4	15
MOTA	2462		56.52	5 10.342 12	8.192 1.00 28.3	35
MOTA	2463	CB TRP A 310	56.94	8.872 12	8.132 1.00 23.9	₹5
ATOM	2464	CG TRP A 310	56.87	8.203 129	9.479 1.00 29.6	
MOTA	2465		55.69		0.263 1.00 31.4	
ATOM-	2466		56.11			
	2467					
MOTA			54.329			10
ATOM	2468	CD1 TRP A 310	57.926			
ATOM	2469	NE1 TRP A 310	57.480		L.436 1.00 30.0	
MOTA	2470	CZ2 TRP A 310	55.213	7.030 132	2.492 1.00 29.9	3
ATOM	2471	CZ3 TRP A 310	53.432	7.831 131	1.062 1.00 29.7	
ATOM	2472	CH2 TRP A 310	53.883	7.259 132	2.265 1.00 24.5	
ATOM	2473	C TRP A 310	57.308			
MOTA	2474	O TRP A 310	56.820			
	2475					
ATOM			58.512		3.984 1.00 29.3	
MOTA	2476	CA CYS A 311	59.305			6
MOTA	2477	CB CYS A 311	60.722	12.538 129	0.479 1.00 30.0	8
MOTA	2478	SG CYS A 311	61.804			.7
MOTA	2479	C CYS A 311	58.612	13.560 130	1.397 1.00 29.2	5
ATOM	2480	O CYS A 311	58.612			
ATOM	2481	N GLU A 312	58.021		.425 1.00 23.1	
ATOM	2482	CA GLU A 312	57:308			
	2483	CB GLU A 312	56.648			
ATOM					.427 1.00 28.9	
MOTA	2484	CG GLU A 312	57.080		.988 1.00 41.6	
MOTA	2485	CD GLU A 312	56.905	_	.059 1.00 44.2	1
ATOM	2486	OE1 GLU A 312	55.813	18.534 129	.658 1.00 54.1	5
ATOM	2487	OE2 GLU A 312	57.860	19.233 129	.290 1.00 43.9	0
ATOM	2488	C GLU A 312	56.204	15.225 130		
ATOM	2489	O GLU A 312	56.120	157.869 131		
ATOM	2490	N LEU A 313	55.343		.388 1.00 31.0	
ATOM	2491	CA LEU A 313	54.231		.266 1.00 36.2	
	2492	CB LEU A 313				_
MOTA			53.337		.604 1.00 28.8	
ATOM	2493	CG LEU A 313	52.493	13.342 129		
ATOM	2494	CD1 LEU A 313	51.818	12.146 128	.788 1.00 33.0	5
MOTA	2495	CD2 LEU A 313	51.471	14.357 129	.914 1.00 27.2	
ATOM	2496	C LEU A 313	54.685	13.377 132		
MOTA	2497	O LEU A 313	54.131	13.730 133		
MOTA	2498	N SER A 314	55.688			
		CA SER A 314		12.508 132		
MOTA	2499		56.233	11.880 133		
ATOM	2500	CB SER A 314	57.183	10.743 133		
MOTA	2501	OG SER A 314	56.517	9.761 132	.628 1.00 45.8	В
ATOM	2502	C SER A 314	57.002	12.846 134	.659 1.00 31.54	4
ATOM	2503	O SER A 314	57.339	12.513 135		
ATOM	2504	N GLY A 315	57.312	14.021 134		
	2505	CA GLY A 315	58.057	14.996 134		
ATOM			59.518			
ATOM	2506					
ATCM	2507	O GLY A 315	60.138	15.049 136		
ATOM	2508	N ARG A 316	60.089	13.862 134	.181 1.00 39.32	3

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MOTA	2509	CA ARG A 316		61.490	13.501 134.332	1.00 39.06
MOTA	2510			61.641		1.00 39.00
MOTA	2511			61.233	11.226 133.184	1.00 39.10
ATOM	2512			61.426	9.744 133.429	
ATOM	2513			60.461	9.744 133.429	
MOTA	2514				9.229 134.389	
ATOM	2519			60.524	8.021 134.926	
				61.511	7.209 134.598	
ATOM	2516			59.583	7.621 135.768	
MOTA	2517			62.369	14.083 133.230	
MOTA	2518			61.910	14.325 132.111	
MOTA	2519			63.633	14.325 133.564	1.00 41.26
ATOM	2520			64.580	14.905 132,619	1.00 44.42
ATOM	2521	= = :	-	65.901	15.249 133.317	1.00 46.84
MOTA	2522			65.756	15.996 134.629	1.00 57.66
ATOM	2523			65.212	15.113 135.743	1.00 65.46
MOTA	2524	OE1 GLU A 317		65.871	14.101 136.073	1.00 68.38
MOTA	2525	OE2 GLU A 317		64.129	15.425 136.287	1.00 67.24
MOTA	2526			64.873	13.962 131.462	1.00 38.45
ATOM	2527	O GLU A 317		64.977	12.748 131.636	1.00 38.84
ATOM	2528	N VAL A 318		65.010	14.525 130.275	1.00 37.64
ATOM	2529			65.315	13.720 129.108	1.00 37.84
ATOM	2530	CB VAL A 318		64.858	14.417 127.810	1.00 39.13
ATOM	2531	CG1 VAL A 318		65.192	13.544 126.610	1.00 42.75
MOTA	2532	CG2 VAL A 318		63.364	14.701 127.867	1.00 41.84
ATOM	2533	C VAL A 318		66.822	13 405 120 027	1.00 42.38
MOTA	2534	0 VAL A 318		67.598	13.495 129.037	1.00 38.45
ATOM	2535	N PRO A 319			14.442 128.910	1.00 36.04
	2536			67.261	12.236 129.156	1.00 39.54
MOTA		CD PRO A 319	-	66.512	10.994 129.397	1.00 40.47
ATOM	2537	CA PRO A 319		68.695	11.949 129.088	1.00 43.85
MOTA	2538	CB PRO A 319	•	68.745	10.439 129.319	1.00 44.12
MOTA	2539	CG PRO A 319		67.419	9.986 128.745	1.00 46.48
ATOM	2540	C PRO A 319		69.228	12.353 127.718	1.00 43.55
MOTA	2541	O PRO A 319		68.563	12.141 126.708	1.00 43.45
MOTA	2542	N GLU A 320		70.420	12.936 127.689	1.00 42.52
ATOM	2543	CA GLU A 320		71.026	13.380 126.440	1.00 45.19
MOTA	2544	CB GLU A 320		72.384	14.032 126.706	1.00 43.86
ATOM	2545	CG GLU A 320		73.121	14.412 125.434	1.00 52.62
MOTA	2546	CD GLU A 320		74.507	14.967 125.697	1.00 52.36
ATOM	2547	OE1 GLU A 320		75.219	15.271 124.720	1.00 56.25
ATOM	. 2548	OE2 GLU A 320		74.883	15.101 126.875	1.00 52.25
MOTA	2549	C GLU A 320		71.223	12.266 125.421	1.00 43.52
MOTA	2550	O GLU A 320		70.876	12.412 124.253	1.00 41.89
ATOM	2551	N LYS A 321		71.781	11.150 125.867	1.00 43.35
MOTA	2552	CA LYS A 321		72.059	10.041 124.969	1.00 43.53
MOTA	2553	CB LYS A 321		73.561	9.808 124.879	1.00 42.78
ATOM	2554	CG LYS A 321		74.238	9.340 126.180	1.00 49.38
ATOM	2555	CD LYS A 321		74.272	10.396 127.307	1.00 57.82
MOTA	2556	CE LYS A 321		72.978	10.497 128.129	1.00 53.81
ATOM	2557	NZ LYS A 321		72.660	9.245 128.883	1.00 54.17
ATOM	2558	C LYS A 321		71.407	8.731 125.345	1.00 41.52
ATOM	2559	O LYS A 321		70.954	8.540 126.469	
MOTA	2560	N LEU A 322		71.378	7.820 124.382	1.00 38.64
ATOM	2561	CA LEU A 322		70.815	6.508 124.613	1.00 40.46
MOTA	2562	CB LEU A 322		70.442	5.845 123.289	1.00 42.22
ATOM	2563	CG LEU A 322		69.595	6.632 122.287	1.00 42.22
ATOM	2564	CD1 LEU A 322		69.204	5.737 121.125	1.00 42.92
ATOM	256.5	CD2 LEU A 322		68.361	7.148 122.967	
ATOM	2566	C LEU A 322		71.918	5.702 125.268	
	2567	O LEU A 322		73.079	5.825 124.884	1.00 41.36
MOTA	2568	N ASN A 323		71.579	4.894 126.265	1.00 44.16
MOTA				72.594		1.00 39.89
MOTA	2569			72.394	4.067 126.895	1.00 40.96
ATOM	2570				3.556 128.259	1.00 43.00
ATOM	2571	CG ASN A 323		70.787	2.886 128.202	1.00 45.59
MOTA	2572	OD1 ASN A 323		70.482	2.151 127.264	1.00 45.71
ATOM	2573	ND2 ASN A 323		69.975	3.114 129.224	1.00 48.08
MOTA	2574	C ASN A 323		72.828	2.894 125.954	1.00 44.88

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	•	
ATO	M 2575 O ASN A 323	72.124 2.739 124 955 1 00 46 43
ATON		7,00 40,41
ATON		73.809 2.062 126.268 1.00 45.98
OTA		74.122 0.938 125.404 1.00 49.82
	11 324	75.386 0.244 125.904 1.00 53.88
ATOM		75.960 -0.711 124.888 1.00 60.09
ATOM	0 077 11914 W 254	75.344 -1.723 124.550 1.00 66.99
MOTA		77.143 -0.386 124.378 1.00 58.06
ATOM	1 2582 C ASN A 324	
ATOM	1 2583 O ASN A.324	
ATOM		
ATOM		1.00 40.33
ATOM		71.106 -1.221 126.318 1.00 46.76
ATOM	525	70.428 -1.328 127.695 1.00 47.65
		71.292 -1.837 128.858 1.00 54 21
ATOM		72.160 -0.750 129.526 1.00 56 87
MOTA	525	73.329 -0.289 128.671 1.00 57.45
ATOM		74.091 0.816 129.307 1.00 58.32
MOTA	. 2591 C LYS A 325	
MOTA	2592 O LYS A 325	
MOTA	2593 N ALA A 326	
MOTA	2594 CA ALA A 326	
ATOM	2595 CB ALA A 326	
ATOM		68.562 2.508 124.616 1.00 42.80
ATOM		69.365 0.940 122.838 1.00 43.80
		68.595 0.625 121.926 1 00 45 17
ATOM	2598 N LYS A 327	70.658 1.191 122 637 1 00 43 46
MOTA	2599 CA LYS A 327	71.235 1.120 121.296 1.00 43.26
ATOM	2600 CB LYS A 327	72.723 1.484 121.311 1.00 44.77
MOTA	2601 CG LYS A 327	73.037 2.892 121.800 1.00 50.87
MOTA	2602 CD LYS A 327	74.544 3.168 121.747 1.00 52.74
MOTA	2603 CE LYS A 327	2.00 32.79
ATOM	2604 NZ LYS A 327	
MOTA	2605 C LYS A 327	
ATOM	2606 O LYS A 327	
ATOM	2607 N GLU A 328	
ATOM	2608 CA GLU A 328	71 276 2 35.93
ATOM	2609 CB GLU A 328	
ATOM	2610 CG GLU A 328	71.875 -3.605 122.135 1.00 43.05
ATOM	2611 CD GLU A 328	73.369 -3.432 122.295 1.00 48.81
ATOM	2612 OE1 GLU A 328	74.096 -3.529 120.963 1.00 52.11
ATOM	2613 OE2 GLU A 328	73.972 -4.574 120.291 1.00 55.44
ATOM	2614 C GLU A 328	74.785 -2.558 120.584 1.00 51.64
ATOM		69.825 -3.030 120.818 1.00 39.51
ATOM	020	69.536 -3.842 119.939 1.00 37.35
		68.911 -2.444 121.582 1.00 36.37
ATOM		67.496 -2.717 121.380 1.00 36.03
ATOM	2618 CB LEU A 329	66.646 -1.958 122.400 1.00 34.66
ATOM	2619 CG LEU A 329	65.133 -2.110 122.213 1.00 33.88
MOTA	2620 CD1 LEU A 329	64.755 -3.572 122.351 1.00 36.21
MOTA	2621 CD2 LEU A 329	64.391 -1.268 123.240 1.00 34.00
MOTA	2622 C LEU A 329	67.120 -2.268 119.971 1.00 33.64
MOTA	2623 O LEU A 329	66.655 -3.061 119.162 1.00 31.29
ATOM	2624 N LEU A 330	67.333 -0.990 119.681 1.00 33.78
MOTA	2625 CA LEU A 330	
MOTA	2626 CB LEU A 330	
MOTA	2627 CG LEU A 330	2.00 30.74
MOTA	2628 CD1 LEU A 330	
ATOM	2629 CD2 LEU A 330	2.00 22.11
ATOM	2630 C LEU A 330	65.028 1.728 118.978 1.00 29.69
		67.729 -1.201 117.246 1.00 36.49
atom atom		67.142 -1.493 116.210 1.00 35.61
		69.005 -1.503 117.455 1.00 37.63
ATOM	2633 CA LYS A 331	69.786 -2.205 116.446 1.00 41.32
	2634 CB LYS A 331	1.256 -2.272 116.874 1.00 44.74
	2635 CG LYS A 331	71.954 -0.919 116.869 1.00 44.68
	2636 CD LYS A 331	73.350 -0.964 117.498 1.00 51.42
MOTA	2637 CE LYS A 331	74.315 -1.889 116.765 1.00 53.71
TOM .	2638 NZ LYS A 331	73.928 -3.327 116.855 1.00 56.15
TOM :	2639 C LYS A 331	69.258 -3.612 116.173 1.00 42.35
	2640 O LYS A 331	1.00 42.33
•	=: 	69.310 -4.086 115.042 1.00 42.68

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```
MOTA
        2641
               N
                   SER A 332
                                    68,734
                                            -4.270 117.200
                                                              1.00 41.56
        2642
                   SER A 332
                                    68.226
                                            -5.629 117.039
                                                              1.00 46.88
ATOM
               CA
                                    68.045
        2643
               CB
                   SER A 332
MOTA
                                            -6.298 118.400
                                                              1.00 42.19
ATOM
        2644
               OG
                   SER A 332
                                    66.959
                                            -5.714
                                                    119.096
                                                              1.00 39.55
ATOM
        2645
               С
                   SER A 332
                                    66.896
                                            -5.687 116.297
                                                              1.00 48.58
                                    66.393
                                            -6.774 116.017
MOTA
        2646
               O
                   SER A 332
                                                              1.00 45.78
MOTA
        2647
               N
                   ILE A 333
                                    66.325
                                            -4.531 115.979
                                                              1.00 48.27
                                            -4.503 115.292
MOTA
        2648
               CA
                   ILE A 333
                                    65.041
                                                              1.00 51.82
                   ILE A 333
                                    64.378
        2649
MOTA
               CB
                                            -3.119 115.402
                                                              1.00 52.16
        2650
               CG2
                   ILE A 333
                                    63.038
                                            -3.122
                                                   114.683
ATOM
                                                              1.00 52.64
        2651
              CG1
                   ILE A 333
                                    64.163
                                            -2.765
                                                   116.871
ATOM
                                                              1.00 52.70
              CD1
                   ILE A 333
                                            -1.402 117.077
MOTA
        2652
                                    63.550
                                                              1.00 56.70
ATOM
        2653
              С
                   ILE A 333
                                    65.112
                                            -4.887
                                                   113.820
                                                              1.00 53.43
MOTA
        2654
              0
                   ILE A 333
                                    66.118
                                            -4.675 113.145
                                                              1.00 56.45
                                    64.016
                                            -5.461 113.344
ATOM
        2655
              N
                   ASP A 334
                                                              1.00 55.53
MOTA
        2656
              CA
                   ASP A 334
                                    63.865
                                            -5.892
                                                   111.962
                                                              1.00 59.58
        2657
                   ASP A 334
                                            -7.040 111.918
              CB
                                   62.845
                                                              1.00 62.69
MOTA
                                   61.546
        2658
                   ASP A 334
MOTA
              CG
                                            -6.712 112.664
                                                              1.00 66.23
                                                              1.00 63.25
ATOM
        2659
              OD1
                  ASP A 334
                                   60.795
                                            -5.814
                                                   112.227
                                            -7.354 113.704
                                                              1.00 63.45
        2660
              OD2
                  ASP A 334
                                   61.277
ATOM
                                   63.385
                   ASP A 334
                                            -4.705 111.125
                                                              1.00 60.81
ATOM
        2661
              C
ATOM
        2662
              0
                   ASP A 334
                                   62.239
                                            -4.673
                                                   110.681
                                                              1.00 59.47
                   PHE A 335
                                   64.266
                                            -3.736
MOTA
        2663
              N
                                                   110.889
                                                              1.00 60.00
              CA
                  PHE A 335
                                   63.864
                                            -2.545
                                                   110.147
                                                              1.00 59.37
ATOM
        2664
                  PHE A 335
                                   64.247
                                            -1.298
ATOM
        2665
              CB
                                                   110.952
                                                              1.00 53.38
                                                             1.00 49.13
ATOM
        2666
              CG
                  PHE A 335
                                   63.895
                                            ~0.013 110.275
MOTA
                  PHE A 335
                                   62.618
       2667
              CD1
                                             0.189
                                                   109.770
                                                              1.00 44.32
                                   64.845
                                             0.993 110.127
       2668
              CD2
                  PHE A 335
                                                              1.00 49.91
ATOM
                                             1.373
                                   62.288
MOTA
       2669
              CE1
                  PHE A 335
                                                   109.122
                                                              1.00 42.74
       2670
              CE2
                  PHE A 335
                                   64.526
                                             2.180
                                                   109.483
ATOM
                                                              1.00 46.40
                                             2.370 108.978
       2671
              CZ
                  PHE A 335
                                   63.244
                                                              1.00 42.77
ATOM
                  PHE A 335
       2672
              C
                                   64.334
                                            -2.399 108.696
ATOM
                                                              1.00 60.85
                                   63.689
                                            -2.914 107.785
MOTA
       2673
              0
                  PHE A 335
                                                              1.00 66.28
       2674
                  GLU A 336
                                   65.430
                                            -1:671 108.493
              N
                                                              1.00 57.40
MOTA
       2675
                  GLU A 336
                                   66.015
                                            -1.411 107.174
                                                              1.00 58.96
ATOM
              CA
                  GLU A 336
                                   65.782
                                            -2.579
MOTA
       2676
              CB
                                                   106.211
                                                              1.00 62.66
       2677
              CG
                  GLU A 336
                                   66.417
                                           -2.377 104.846
MOTA
                                                              1.00 68.51
       2678
              CD
                  GLU A 336
                                   66.277
                                            -3.590
                                                   103.943
                                                              1.00 73.21
ATOM
                                   66.753
                                            -4.678
                  GLU A 336
                                                   104.333
MOTA
       2679
              OE1
                                                             1.00
                                                                   73.30
                                                              1.00 75.74
       2680
              OE2
                  GLU A 336
                                   65.697
                                            -3.457
                                                   102.843
ATOM
ATOM
       2681
              С
                  GLU A 336
                                   65.460
                                            -0.124
                                                   106.576
                                                             1.00 55.70
                                           -0.023
                                                   106.253
              o
                  GLU A 336
                                   64.281
       2682
                                                             1.00 55.28
MOTA
                                                   106.432
                  GLU A 337
                                            0.857
MOTA
       2683
             N
                                   66.338
                                                             1.00 54.75
MOTA
       2684
             CA
                  GLU A 337
                                   65.986
                                            2.167
                                                   105.905
                                                             1.00 55.99
                                   67.221
                                            3.065
                                                             1.00 51.75
       2685
             CB
                  GLU A 337
                                                   105.983
MOTA
                                            4.536
                                                   106.092
                                   66.926
                                                             3 00 52.28
       2686
                  GLU A 337
ATOM
             CG
MOTA
       2687
             CD
                  GLU A 337
                                   66.184
                                            4.891
                                                   107.366
                                                               .00 43.72
                                   66.705
                                            4.640 108.474
                                                             3 00 42.21
       2688
             OE1
                 GLU A 337
ATOM
                                   65.072
                                            5.425 107.256
                 GLU A 337
             OE2
       2689
                                                             1.00 47.31
ATOM
       2690
             C
                  GLU A 337
                                   65.485
                                            2.064
                                                   104.460
                                                             1.00
                                                                   57.56
ATOM
                                   66.087
                                            1.377 103.639
       2691
             0
                  GLU A 337
                                                             1.00 58.29
ATOM
                                            2.745 104.151
2.710 102.805
                                   64.385
       2692
             N
                  PHE A 338
                                                             1.00 60.26
MOTA
       2693
             CA
                 PHE
                      A 338
                                   63.814
                                                             1.00
                                                                  61.69
ATOM
                  PHE A 338
                                  62.561
                                            3.582 102.723
       2694
             CB
                                                             1.00 60.86
MOTA
                                            3.493 101.401
                                  61.845
                                                             1.00 61.92
MOTA
       2695
             CG
                 PHE A 338
                                                   101.094
       2696
             CD1
                 PHE A 338
                                  61.054
                                            2.391
                                                             1.00 62.99
ATOM
       2697
             CD2
                 PHE A 338
                                  61.970
                                            4.508
                                                   100.458
                                                             1.00 62.74
ATOM
                                  60.392
                                            2.302
                                                    99.868
             CE1
                                                             1.00 65.11
ATOM
       2698
                 PHE
                     A 338
                                                    99.228
       2699
             CE2
                 PHE
                     Α
                        338
                                  61.315
                                            4.428
                                                             1.00 64.62
MOTA
                                  60.523
                                            3.322
                                                    98.934
       2700
                 PHE
                     A 338
                                                             1.00 63.87
             CZ
ATOM
                                            3.208
                                  64.818
                                                   101.773
                                                             1.00 64.33
       2701
             C
                 PHE
                     A 338
ATOM
       2702
                 PHE
                     A
                        338
                                  64.803
                                            2.781
                                                   100.616
                                                             1.00 62.45
             0
ATOM
                                            4.130
                                                   102.194
                                  65.677
                 ASP
                     A 339
                                                             1.00 64.11
       2703
             N
ATOM
                                            4.684 101.310
                                  66.689
                 ASP A 339
       2704
             CA
                                                             1.00 67.42
ATOM
                                            6.206 101.248
       2705
                 ASP
                     A 339
                                  66.565
                                                             1.00 67.20
             CB
ATCM
                                  67.647
                                            6.838 100.402
                 ASP
                                                             1.00 68.03
       2706
             CG
ATOM
```

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3.000	2707	OD1 ASP A 339	CD	206 6	430 00 00	
MOTA					432 99.22	
MOTA	2708	OD2 ASP A 339	68	.346 .7.	740 100.904	4 1.00 64.15
MOTA	2709	C ASP A 339	68	.088 4.	295 101.763	
ATOM	2710					
					628 102.869	
MOTA	2711	N ASP A 340	68	.796 3.	588 100.888	3 1.00 71.90
ATOM	2712	CA ASP A 340	70	.151 3	111 101.149	
ATOM	2713	CB ASP A 340				
					601 99.848	
MOTA	2714	CG ASP A 340	69.	.953 1.	511 99.199	5 1.00 76.03
ATOM	2715	OD1 ASP A 340	69	761 0.	449 99.823	3 1.00 76.90
	2716	OD2 ASP A 340				1.00 76.90
MOTA					718 98.05	
MOTA	2717	C ASP A 340	- 71.	069 4	155 101.76 <i>6</i>	1.00 72.77
ATOM	2718	O ASP A 340	71	618 3.	946 102.845	
	2719		71.			1.00 73.06
MOTA					275 101.074	
MOTA	2720	CA GLU A 341	72.	112 6.3	341 101.557	1.00 74.56
ATOM	2721	CB GLU A 341	72	917 6.5	924 100.390	
	2722					1.00 //.00
ATOM					034 100.792	
ATOM	2723	CD GLU A 341	74.	924 7.5	571 101.794	1.00 85.34
ATOM	2724	OE1 GLU A 341	75.	718 6.6	569 101.450	
	2725					1.00 00.04
MOTA					106 102.924	
MOTA	. 2726	C GLU A 341	71.	327 7.4	153 102.245	1.00 72.47
ATOM	. 2727	O GLU A 341	70.	822 8.3	364 101.589	1.00 76.75
	2728	N VAL A 342		220 7 3	01 102.505	
MOTA					881 103.566	
MOTA	2729	CA VAL A 342	70.	503 8.3	393 104.323	1.00 64.84
ATOM	2730	CB VAL A 342	69.	160 7.8	353 104.850	
ATOM	2731	CG1 VAL A 342	68.		194 103.701	
		CGI VAL A J42				
MOTA	2732	CG2 VAL A 342	69.		37 105.722	1.00 65.37
MOTA	2733	C VAL A 342	71.	305 8.8	371 105.520	1.00 61.75
ATOM	2734	O VAL A 342	71.		66 105.795	
MOTA	2735	N ASP A 343	71.		25 106.225	1.00 56.79
ATOM-	2736	CA ASP A 343	72.	692 8.2	29 107.417	1.00 54.53
ATOM	2737	CB ASP A 343	73.		40 107.158	
	2738					
MOTA		CG ASP A 343	74.		60 108.388	
ATOM	2739	OD1 ASP A 343	75.	298 10.6	44 108.357	1.00 65.36
MOTA	2740	OD2 ASP A 343	74.		18 109.387	
	2741					
ATOM			71.		75 108.534	1.00 50.70
ATOM	2742	O ASP A 343	71.	142 9.8	59 108.651	1.00 46.00
ATOM	2743	N ARG A 344	71.:	328 7.7	17 109.341	1.00 46.20
	2744	CA ARG A 344	70.4			
ATOM					04 110.463	1.00 41.18
MOTA	2745	CB ARG A 344	69.3		68 110.299	1.00 39.81
MOTA	2746	CG ARG A 344	68.2	289 7.7	11 109.098	1.00 35.08
ATOM	2747	CD ARG A 344	68.0			1.00 28.37
ATOM	2748	NE ARG A 344	67.1		45 108.036	1.00 30.90
ATOM	2749	CZ ARG A 344	67.0	013 10.9	09 107.649	1.00 31.05
MOTA	2750	NH1 ARG A 344	67.6		74 108.258	1.00 30.49
	2751	NH2 ARG A 344				
MOTA			66.2		12 106.646	1.00 31.76
ATOM	2752	C ARG A 344	71.1	.47 7.5	61 111,742	1.00 38.46
MOTA	2753	O ARG A 344	70.5	16 7.3	70 112.773	1.00 34.99
	2754	N SER A 345	72.4			1 00 33 05
MOTA					18 111.662	1.00 33.97
MOTA	2755	CA SER A 345	73.2	61 6.9	81 112.795	1.00 33.68
MOTA	2756	CB SER A 345	74.7	42 6.9	72 112.404	1.00 39.11
	2757	OG SER A 345	75.1		60 111.990	1.00 42.80
ATOM						
ATOM		C SER A 345	73.0	54 7.8	26 114.053	1.00 31.83
ATOM	2759	O SER A 345	73.1	00 7.33	14 115.167	1.00 24.35
	2760	N TYR A 346	72.8		19 113.877	1.00 33.10
MOTA						1.00 33.10
MOTA		CA TYR A 346	72.6		03 115.015	1.00 34.50
MOTA	2762	CB TYR A 346	72.3	97 11.43	37 114.522	1.00 35.16
ATOM		CG TYR A 346	71.1		15 113.659	1.00 39.69
						1 00 22 22
ATOM	-	CD1 TYR A 346	69.9		14 114.227	1.00 36.57
MOTA	2765	CEl TYR A 346	68.7		10 113.424	1.00 40.23
ATOM	2766	CD2 TYR A 346	71.2		4 112.270	1.00 39.04
		CE2 TYR A 346	70.1			1 00 30 60
ATOM					7 111.463	1.00 38.65
MOTA	2768	CZ TYR A 346	68.8	9U 11.86	4 112.041	1.00 37.64
MOTA	2769	OH TYR A 346	67.7	76 11.98	2 111.234	1.00 32.48
		C TYR A 346	71.4		0 115.874	1.00 37.72
ATOM	-					1.00 37.72
ATCM	-	O TYR A 346.	71.3		9 117.074	1.00 35.48
ATCM	2772	N MET A 347	70.4	72 8.86	9 115.265	1.00 35.36

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```
ATOM
         2773
                   MET A 347
                                    69.295
               CA
                                              8.418 116.004
                                                              1.00 36.97
MOTA
         2774
                   MET A 347
                                    68.226
                                              7.868 115.052
               CB
                                                              1.00 33.45
         2775
                   MET A 347
MOTA
                                    67.853
               CG
                                              8.809 113.921
                                                              1.00 28.09
         2776
MOTA
               SD
                   MET A 347
                                    66.471
                                             8.194 112.943
                                                              1.00 32,14
        2777
                                    67.058
MOTA
               CE
                   MET A 347
                                              6.647 112.457
                                                              1.00 39.25
        2778
                                   69.632
ATOM-
               C
                   MET A 347
                                              7.363 117.055
                                                              1.00 34.67
        2779
                                    68.890
MOTA
               0
                   MET A 347
                                             7.187 118.016
                                                              1.00 33.46
        2780
                   LEU A 348
                                   70.747
MOTA
               N
                                             6.663 116.880
                                                              1.00 36.42
        2781
               CA
                   LEU A 348
                                   71.137
MOTA
                                             5.641 117.846
                                                              1.00
                                                                  34.57
MOTA
        2782
               CB
                   LEU A 348
                                   71.841
                                              4.476 117.152
                                                              1.00 34.16
        2783
MOTA
               CG
                   LEU A 348
                                   71.066
                                             3.655 116.121
                                                              1.00 35.61
        2784
               CDI
                   LEU A 348
                                   72.010
ATOM
                                             2.660 115.469
                                                              1.00 36.58
ATOM
        2785
               CD2
                   LEU A 348
                                   69.906
                                             2.935 116.788
                                                              1.00 42.34
        2786
                   LEU A 348
                                   72.075
MOTA
               С
                                             6.232 118.885
                                                              1.00 37.10
        2787
               0
                   LEU A 348
                                   72.583
ATOM
                                             5.513 119.745
                                                              1.00 37.75
MOTA
        2788
               N
                   GLU A 349
                                   72.295
                                             7.541 118.801
                                                              1.00 38.31
        2789
                   GLU A 349
MOTA
               CA
                                   73.192
                                             8.231 119.726
                                                              1.00 42.86
        2790
               CB
                   GLU A 349
                                   74.150
ATOM
                                             9.136 118.948
                                                             1.00 47.04
ATOM
        2791
               CG
                   GLU A 349
                                   74.942
                                             8.426 117.867
                                                             1.00 53.27
        2792
                   GLU A 349
                                   75.828
                                             7.325 118.410
MOTA
               CD
                                                             1.00 58.36
        2793
              OE1
                  GLU A 349
                                   76.681
                                             7.619
MOTA
                                                   119.279
                                                             1.00 59.59
        2794
                                   75.669
ATOM
              OE2
                   GLU A 349
                                             6.166 117.963
                                                             1.00 59.17
        2795
                   GLU A 349
                                   72.458
                                                             1.00 41.72
ATOM
                                             9.080 120.752
        2796
ATOM
              0
                   GLU A 349
                                   72.564
                                             8.852 121.959
                                                             1.00 42.27
        2797
                   THR A 350
MOTA
              N
                                   71.716
                                            10.067 120.261
                                                             1.00 37.05
        2798
                   THR A 350
MOTA
              CA
                                   70.992
                                            10.976 121.135
                                                             1.00 40.78
MOTA
        2799
              CB
                   THR A 350
                                   71.468
                                            12.418 120.900
                                                             1.00 41.69
MOTA
        2800
              OG1
                                   71.359
                  THR A 350
                                            12.733 119.508
                                                             1.00 43.25
        2801
                  THR A 350
MOTA
              CG2
                                   72.918
                                            12.575 121.325
                                                             1.00 44.11
MOTA
        2802
              C
                   THR A 350
                                   69.474
                                            10.911 120.968
                                                             1.00 41.47
        2803
              0
                   THR A 350
                                   68.968
                                            10.608 119.884
                                                             1.00 39.72
ATOM
                                   68.760
        2804
              N
                  LEU A 351
                                           11.199 122.054
ATOM
                                                             1.00 38.68
        2805
                                   67.299
MOTA
              CA
                  LEU A 351
                                           11.185 122.056
                                                             1.00 35.91
MOTA
        2806
              CB
                  LEU A 351
                                   66.763
                                           11:210 123.487
                                                             1.00 35.94
                                   66.752
        2807
              CG
                  LEU A 351
                                            9.890 124.251
ATOM
                                                             1.00 37.67
       2808
              CD1
                  LEU A 351
                                   66.290
ATOM
                                           10.118 125.677
                                                             1.00 40.11
ATOM
       2809
              CD2
                  LEU A 351
                                   65.813
                                            8.920 123.544
                                                             1.00 39.29
       2810
              С
                  LEU A 351
                                   66.679
                                           12.342 121.294
ATOM
                                                             1.00 37.76
                                   65.747
       2811
                  LEU A 351
ATOM
              0
                                           12.149 120.512
                                                             1.00 34.86
ATOM
       2812
              N
                  LYS A 352
                                   67.192
                                           13.544 121.525
                                                             1.00 34.79
MOTA
       2813
              CA
                  LYS A 352
                                   66.651
                                           14.724 120.870
                                                             1.00 38.73
                                  66.676
66.062
       2814
              CB
                  LYS A 352
                                           15.911 121.835
                                                             1.00 36.48
ATOM
       2815
                  LYS A 352
MOTA
              CG
                                           15.580 123.179
                                                             1.00 42.08
MOTA
       2816
              CD
                  LYS A 352
                                  66.202
                                           16.701 124.196
                                                             1.00 43.22
ATOM
       2817
              CE
                  LYS A 352
                                  65.349
                                           17.901 123.845
                                                             1.00 49.81
       2818
                                  65.342
MOTA
             NZ
                  LYS A 352
                                           18.880 124.972
                                                             1.00 52.70
                  LYS A 352
ATOM
       2819
              C
                                  67.425
                                           15.063 119.610
                                                             1.00 38.77
       2820
              0
                  LYS A 352
                                  68.654
                                           15.098 119.601
                                                             1.00 36.21
MOTA
                  ASP A 353
       2821
                                  66.697
MOTA
             N
                                           15.293 118.530
                                                             1.00 37.69
       2822
                  ASP A 353
                                                             1.00 39.89
MOTA
             CA
                                  67.337
                                           15.647 117.286
       2823
             CB
                  ASP A 353
                                  66.532
                                                             1.00 43.53
ATOM
                                           15.075 116.110
       2824
             CG
                  ASP A 353
                                  65.058
                                           15.368 116.211
MOTA
                                                             1.00 47.99
       2825
             OD1
                  ASP A 353
MOTA
                                  64.253
                                           14.623 115.593
                                                             1.00 38.06
                                  64.706
       2826
             OD2
                  ASP A 353
                                           16.352 116.898
MOTA
                                                             1.00 52.94
       2827
                  ASP A 353
MOTA
             C
                                  67.457
                                           17.165 117.247
                                                             1.00 39.20
       2828
                  ASP A 353
MOTA
             0
                                  66.890
                                           17.861 118.092
                                                             1.00 35.66
       2829
             N
                  PRO A 354
                                  68.244
                                           17.696 116.302
MOTA
                                                             1.00 40.75
                                  69.047
       2830
             CD
                  PRO A 354
                                           17.005 115.279
                                                             1.00 40.05
ATOM
                  PRO A 354
                                  68.426
MOTA
       2831
             CA
                                           19.145 116.179
                                                            1.00 38.41
       2832
             СB
                  PRO A 354
                                  69.534
                                           19.250 115.140
                                                            1.00 36.24
MOTA
                                           18.095 114.225
      2833
             CG
                  PRO A 354
                                  69.190
                                                            1.00 39.41
ATOM
      2834
             C
                  PRO A 354
                                  67.144
                                           19.780 115.689
                                                            1.00 37.28
MOTA
      2835
                  PRO A 354
                                  66.299
             0
                                           19.106 115.094
                                                            1.00 31.87
MOTA
      2836
             N
                  TRP A 355
                                  66.993
                                           21.074 115.934
                                                            1.00 37.87
MOTA
                                  65.804
      2837
             CA
                 TRP A 355
                                           21.757 115.472
                                                            1.00 40.04
ATOM
                                           23.157 116.080
      2838
                  TRP A 355
                                  65.714
                                                            1.00 42.85
MOTA
             CB
```

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MOTA	2839	CG TRP A 355	64.333	23.471 116.569	1.00 53.01
MOTA	2840	CD2 TRP A 355	63.439	24.463 116.047	1.00 54.23
MOTA	2841	. CE2 TRP A 355	62.234		1.00 52.59
MOTA	2842	CE3 TRP A 355	63.538	25.422 115.028	1.00 52.82
MOTA	2843	CD1 TRP A 355	63.655		1.00 51.80
	•				
MOTA	2844	NE1 TRP A 355	62.393	23.368 117.708	1.00 51.38
ATOM	2845	CZ2 TRP A 355	61.134	25.197 116.532	1.00 51.28
MOTA	2846		62.444		1.00 53.22
ATOM	2847	CH2 TRP A 355	61.257	26.126 115.531	1.00 50.50
ATOM	2848		65.935	21.836 113.954	1.00 37.34
MOTA	2849	O TRP A 355	67.041	21.929 113.422	1.00 39.01
MOTA	2850	N ARG A 356	64.809	21.764 113.259	1.00 36.31
ATOM	2851	CA ARG A 356	64.797	21.820 111.802	1.00 35.22
MOTA	2852	CB ARG A 356	64.317	20.469 111.252	1.00 33.35
MOTA	2853	CG ARG A 356	65.310	19.340 111.564	1.00 34.50
	2854	CD ARG A 356	64.729	17.927 111.468	1.00 28.06
MOTA					
MOTA	2855	NE ARG A 356	65.745	16.956 111.870	1.00 24.79
MOTA	2856	CZ ARG A 356	65.499	15.703 112.236	1.00 27.56
	2857	NH1 ARG A 356	64.253	15.237 112.259	
MOTA					1.00 19.09
ATOM	2858	NH2 ARG A 356	66.502	14.918 112.604	1.00 21.34
ATOM	2859	C ARG A 356	63.874	22.955 111.390	1.00 33.74
	2860				1.00 34.68
MOTA			62.746		
MOTA	2861	N GLY A 357	64.361	24.181 111.550	1.00 35.60
MOTA	2862	CA GLY A 357	63.556	25.345 111.220	1.00 35.43
	2863	C GLY A 357	63.719	25.932 109.830	1.00 38.08
MOTA					
MOTA	2864	O GLY A 357	64.112	25.250 108.885	1.00 37.29
ATOM	2865	N GLY A 358	63.406	27.218 109.721	1.00 39.67
	2866	CA GLY A 358	63.493	27.925 108.457	1.00 36.36
MOTA					
MOTA	2867	C GLY A 358	62.398	28.966 108.499	1.00 39.45
ATOM	2868	O GLY A 358	61.763	29.131 109.539	1.00 37.58
	2869	N GLU A 359	62.163	29.662 107.391	1.00 40.89
ATOM					
ATOM	2870	CA GLU A 359	61.121	30.682 107.358	1.00 41.37
ATOM	2871	CB GLU A 359	61.310	31.627 106.172	1.00 44.64
	2872	CG GLU A 359	60.956	30.977 104.848	1.00 52.13
ATOM					
MOTA	. 2873	CD GLU A 359	60.833	31.973 103.708	1.00 59.14
MOTA	2874	OE1 GLU A 359	60.448	31.551 102 <i>.</i> 593	1.00 60.47
ATOM	2875	OE2 GLU A 359	61.119	33.173 103.923	1.00 57.77
MOTA	2876	C GLU A 359	59.770	30.006 107.200	1.00 38.02
ATOM	2877	O GLU A 359	59.689	28.828 106.850	1.00 35.29
ATOM	2878	N VAL A 360	58.708	30.762 107.441	1.00 36.81
	-			30.237 107.291	
MOTA	2879	CA VAL A 360	57.363		1.00 35.97
ATOM	2880	CB VAL A 360	56.401	30.789 108.368	1.00 34.90
MOTA	2881	CG1 VAL A 360	54.999	30.251 108.133	1.00 36.53
		CG2 VAL A 360	56.888	30.393 109.755	1.00 37.06
ATOM	2882				
ATOM	2883	C VAL A 360	56.886	30.690 105.928	1.00 36.74
ATOM	2884	O . NAT W 360	56.661	31.881 105.712	1.00 34.90
	2885	N ARG A 361	56.753	29.741 105.004	1.00 35.48
ATOM					
ATOM	2886	CA ARG A 361	56.301	30.049 103.652	1.00 38.21
ATOM	2887	CB ARG A 361	56.152	28.776 102.815	1.00 39.76
	2888	CG ARG A 361	57.416	28.342 102.098	1.00 39.93
ATOM					
ATOM	2889	CD ARG A 361	57.225	26.963 101.486	1.00 38.68
ATCM	2890	NE ARG A 361	57.112	25.940 102.525	1.00 39.72
	2891	CZ ARG A 361	56.952	24.643 102.286	1.00 38.79
ATOM					
ATOM	2892	NH1 ARG A 361	56.881	24.200 101.036	1.00 32.40
MOTA	2893	NH2 ARG A 361	56.899	23.785 103.297	1.00 36.58
	2894	C ARG A 361	54.996	30.807 103.603	1.00 38.98
ATOM				30.636 104.452	1.00 39.07
MOTA	2895	O , ARG A 361	54.120		
ATCM	2896	N LYS A 362	54.880	31.634 102.573	1.00 39.95
	2897	CA LYS A 362	53.709	32.459 102.339	1.00 42.73
MOTA					
ATCM	2898	CB LYS A 362	53.931		1.00 44.92
ATOM	2899	CG LYS A 362	54.995	34.390 101.219	1.00 55.45
	2900	CD LYS A 362	56.351	33.842 101.671	1.00 58.28
ATCM			56.907		1.00 57.04
ATCM	2901	CE LYS A 362			1.00 5/.04
ATOM	2902	NZ LYS A 362	58.224	32.283 101.151	1.00 58.75
	2903	C LYS A 362	52.434	31.634 102.200	1.00 40.51
ATOM			51.391	31.996 102.748	1.00 36.10
ATCM	2904	O LYS A 362	コエ・コユエ	JA. 330 AV2. /48	1.00 30.10

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ATOM	290	N GLUA	363	52.506	30.527	7 101.469	1.00	37.79
ATOM	2900	CA GLUA	363	51.313	29.70			40.96
ATOM	290			51.587	28.530			43.62
ATOM	2908			52.729		100.739		47.01
ATOM	2909			52.995	26.547			51.65
ATOM	2910			52.080	25.737			45.63
MOTA	2911			54.116	26.526			48.44
MOTA	2912			50.788		102.636	1 00	37.74
ATOM	2913			49.582	29.113			34.79
ATOM	2914			51.691		103.564		33.64
ATOM	2915			51.274		104.886	1.00	32.39
ATOM	2916			52.484		105.749	1.00	33.99
ATOM	2917			52.018		107.160	1.00	36.90
ATOM	2918		364	53.198		107.100		29.56
ATOM	2919			50.506		105.589		34.33
ATOM	2920			49.454		105.389		29.80
ATOM	2921			51.027		105.499		38.12
ATOM	2922		65	50.381		106.119		36.77
ATOM	2923			51.255	33.204			
ATOM	2924	CG LYS A 3		52.629	33.204			37.98
ATOM	2925	CD LYS A 3		53.449	34.357			37.99
ATOM	2926	CE LYS A 3	65	54.837		100.429		35.50
ATOM	2927	NZ LYS A 3		55.674		106.877		40.35
ATOM	2928	C LYS A 3		49.025		105.468		43.74
ATOM	2929	O LYS A 3		48.038		105.468		36.62
ATOM	2930	N ASP A 3		48.968		104.147		33.53
ATOM	2931	CA ASP A 3	66	47.708		104.147	1.00	37.05
ATOM	2932	CB ASP A 3		47.906		101.929		37.72
ATOM	2933	CG ASP A 3		48.833		101.427		40.57
ATOM	2934	OD1 ASP A 3		19.078		102.176		43.98
ATOM	2935	OD2 ASP A 3	66	19.297		100.269		41.96
ATOM	2936	C ASP A 3		16.670		103.862		39.24
ATOM	2937	O ASP A 3		15.497		104.029	1.00	39.04
ATOM	2938	N THR A 3		17.096		104.031		38.99
ATOM	2939	CA THR A 3		6.167	28.935			36.80
ATOM	2940	CB THR A 3		6.868	27.560		1 00	33.84
ATOM	2941	OG1 THR A 3		7.332	27.167		1.00	
ATOM	2942	CG2 THR A 3	_	5.904	26.509		1.00	
ATOM	2943	C THR A 3		5.532	29.257		1.00	
ATOM	2944	O THR A 3		4.307		105.931	1.00	
ATOM	2945	N LEU A 3	_	6.363		106.776	1.00	
ATOM	2946	CA LEU A 36		5.850		108.095	1.00	
ATOM	2947	CB LEU A 36		6.997		109.077	1.00	
ATOM	2948	CG LEU A 36	58 4	7.545		109.794	1.00	
ATOM	2949	CD1 LEU A 36	58 4	6.449		110.688		37.35
MOTA	2950	CD2 LEU A 36	8 4	8.014		108.797	1.00	
ATOM	2951	·C LEU A 36	58 4	4.957		107.994		38.12
MOTA	2952	O LEU A 36	8 4	3.968		108.719	1.00	
ATOM	2953	N GLU A 36	9 4	5.307	32.063	107.086	1.00	
ATOM	2954	CA GLU A 36	9 4	4.509		106.866	1.00	
ATOM	2955	CB GLU A 36	9 4:	5.128	34.126	105.765	1.00	47.38
ATOM	2956	CG GLU A 36		6.020		106.283	1.00 9	
ATOM	2957	CD GLU A 36	9 4:	5.227	36.306	106.996	1.00	
ATOM	2958	OE1 GLU A 36	9 4	5.846		107.526	1.00	
ATOM	2959	OE2 GLU A 36	9 4:	3.980		107.016	1.00	
ATOM	2960	C GLU A 36				106.466	1.00 4	
ATOM	29.61	O GLU A 36	9 42			107.095	1.00 4	
MOTA	2962	N LYS A 37	0 42			105.417	1.00 4	
ATOM	2963	CA LYS A 37				104.977	1.00 4	
ATOM	2964	CB LYS A 37	0 41			103.704	1.00 4	
ATOM	2965	CG LYS A 37	0 42		31.546	102.419	1.00 4	
ATOM	2966	CD LYS A 37			31.514	102.019	1.00 5	
MOTA	2967	CE LYS A 37			30.116	101.561	1.00 5	
ATOM	2968	NZ LYS A 37		5.339		101.091	1.00 5	
ATOM	2969	C LYS A 37				106.069	1.00 4	
MOTA	2970	O LYS A 37	0 39	7.745	30.977	106.248	1.00 4	1.34

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MOTA	297:	l N	AL	A A 371	41.715	30 037	106.802	1.00 39.56
ATOM	2972	2 C		A A 371				
			-		41.120		107.861	
MOTA	2973			A A 371	42.193	28.440	108.594	1.00 39.11
ATOM	2974	4 C	AL	A A 371	40.365	30.132	108.837	
MOTA	2975	5 0	AT.	A A 371	39.230	29.829		
	2976							
MOTA				S A 372	40.981	31.239	109.227	1.00 46.62
MOTA	297.7	7 C	7 LTA	S A 372	40.391	32.178	110.192	1.00 48.18
MOTA	2978	3 C	LYS	S A 372	39.052	32.698		
ATOM	2979			S A 372	38.294			
						33.318		
MOTA	2980			S A 372	41.334	33.364	110.413	
MOTA	. 2981	. CG	LYS	S A 372	42.804	32.949	110.510	
ATOM	2982	CI	1.YS	5 A 372	43.746		110.752	1.00 20.00
	2983							
MOTA				A 372	45.216		110.849	1.00 20.00
MOTA	2984	NZ		a 372	46.121	34.850	111.081	1.00 20.00
ATOM	2985	N	ALA	A 373	38.751	32.476	108.397	1.00 57.71
ATOM	2986	CA		A 373	37.492	32.933	107.806	1.00 58.67
	2987			A 373				
ATOM					37.758		106.480	1.00 57.19
MOTA	2988	С		A 373	36.524	31.773	107.594	1.00 59.58
MOTA	2989	0	ALA	A 373	35.432	31.797	108.205	1.00 60.89
ATOM	2990	OX		A 373	36.870	30.853	106.822	1.00 60.10
	3014	CB						
ATOM					54.881	-4.431	56.836	1.00 55.77
MOTA	3015	С	ALA		53.960	-2.137	56.480	1.00 57.58
MOTA	3016	0	ALA	9 2	54.920	-1.720	57.131	1.00 56.75
ATOM	3017	N	ALA		54.263	-3.672		
	3018	CA		- ÷			54.557	1.00 58.22
ATOM			ALA		53.914	-3.584	56.008	1.00 58.47
MOTA	3019	N.	LYS		52.919	~1.376	56.151	1.00 52.79
MOTA	3020	CA	LYS	в 3	52.855	0.022	56.543	1.00 49.68
ATOM	3021	CB	LYS		51.643	0.700		
							55.896	1.00 53.14
MOTA	3022	CG	LYS		51.751	0.785	54.377	1.00 53.37
ATOM	3023	CD	LYS		50.685	1.681	53.786	1.00 55.40
MOTA	3024	CE	LYS	B 3	50.808	1.783	52.277	1.00 59.51
ATOM	3025	NZ	LYS		52.140	2.323	51.884	
								1.00 56.88
MOTA	3026	Ç	LYS		52.849	0.238	58.059	1.00 46.83
ATOM	3027	0	LYS	в 3.	52.389	-0.607	58.830	1.00 41.63
ATOM	3028	N	VAL	в 4	53.376	1.385	58.467	1.00 41.46
ATOM	3029	CA	VAL		53.483		59.871	
						1.751		1.00 40.85
ATOM	3030	CB	VAL		54.893	2.288	60.163	1.00 39.55
ATOM	3031	CG1	. VAL	B 4	55.070	2.541	61.648	1.00 41.23
ATOM	3032	CG2	VAL	B 4	55.916	1.306	59.652	1.00 38.96
MOTA	3033	C	VAL		52.451	2.813		
							60.230	1.00 38.92
ATOM	3034	0	VAL		52.472	3.916	59.691	1.00 42.80
MOTA	3035	N	LYS	в 5	51.559	2.479	61.157	1.00 34.90
ATOM	3036	CA	LYS	B 5	50.501	3.396	61.558	1.00 31.22
ATOM	3037	CB	LYS	в 5	49.133	2.796	61.215	
								1.00 33.76
ATOM	3038	CG	LYS		48.841	2.623	59.726	1.00 36.60
ATOM	3039	CD	LYS		48.667	3.964	59.032	1.00 41.48
ATOM	3040	CE	LYS	3 5	48.234	3.803	57.577	1.00 43.62
ATOM	3041	NZ	LYS		49.215	3.025	56.781	1.00 42.53
	3042							
ATOM		C	LYS		50.512	3.749	63.038	1.00 32.67
ATOM	3043	0	LYS	B 5	51.012	2.995	63.878	1.00 25.78
MOTA	3044	N	LEU :	B 6	49.937	4.906	63.343	1.00 27.07
ATOM	3045	CA	LEU :		49.821	5.379	64.712	1.00 31.09
					49.021			1.00 31.09
MOTA	3046	CB	LEŲ I		50.596	6.696	64.896	1.00 30.13
ATOM	3047	CG	LEU I	B 6	50.691	7.340	66.285	1.00 28.09
ATOM	3048	CD1	LEU I	В 6	49.333	7.827	66.728	1.00 38.87
ATOM	3049		LEU I		51.248		67.282	
						6.338		1.00 24.87
MOTA	3050	C	LEU I		48.324	5.594	64.924	1.00 29.52
ATOM	3051.	0	LEU S	36	47.669	6.287	64.149	1.00 33.36
MOTA	3052	N	ILE E		47.777	4.975	65.960	1.00 28.02
	3053							
MOTA		CA	ILE S		46.361	5.111	66.250	1.00 23.83
MOTA	3054	CB	ILE E		45.736	3.761	66.670	1.00 25.11
MOTA	3055	CG2	ILE E	3 7	44.309	3.974	67.127	1.00 23.59
ATOM	3056	CG1	ILE E		45.690	2.794	65.477	1.00 31.00
	3057	CD1						
MOTA			ILE E		47.021	2.406	64.906	1.00 38.60
ATOM	3058	C	ILE E		46.179	6.130	67.363	1.00 26.49
MOTA	3059	0	ILE E	37	46.766	6.005	68.430	1.00 26.68
					-			

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	MOTA	3060	N GI	Y B	8		45.37	72 7 11	-1 62 16		
	ATOM	3061		ΥB	8				· · · · · · · · · · · · · · · · · · ·		
	ATOM			ΥB	8		45.15				
	ATOM	3063		Ϋ́В			44.21				9
	ATOM	3064		R B	8		43.62			0 1.00 19.7	0
	ATOM-				9		44.08		68.50	9 1.00 26.4	6
	ATOM	3066		RB	9		43.23				
		_		RВ	9		41.74				4
	MOTA	3067		R B	9		40.95	9 12.25	3 68.21	8 1.00 30.35	5
	MOTA	3068		R B	9		41.43			7 1.00 33.42	,
	ATOM	3069		R B	9		43.47	9 12.49	6 69.30	2 1.00 33.42	,
	ATOM	3070		RВ	9		43.88	4 12.17		6 1.00 30.46	:
	MOTA	3071	N LE	UB.	10		43.22			1 1.00 32.05	•
	MOTA	3072	CA LE	UB	10	•	43.39		0 69.91		
	MOTA	3073	CB LE	JВ	10		43.38				
	ATOM	3074	CG LE	JB	10		44.60				
	ATOM	3075	CD1 LET	JВ	10		44.96				
	ATOM	3076	CD2 LEG		10		44.31				
	ATOM	30 7 7	C LE		10		42.27				
	ATOM	3078	O LE		10		42.41				
	ATOM		N ASE		11						
	MOTA		CA ASE		11		41.158				
	ATOM		CB ASE		11		40.013				
	ATOM		CG ASP				38.928				
	ATOM		OD1 ASP		11		38.372				
	ATOM				11		38.013		69.525		
	ATOM		OD2 ASP		11		38.281				
	ATOM		C ASP		11		40.358			1.00 32.19	
			O ASP		11		39.688			1.00 23.44	
	ATOM		N TYR		12		41.386			1.00 28.02	
	ATOM	_	CA TYR		12		41.770				
	ATOM		CB TYR		12		43.011			1.00 28.67	
	MOTA		G TYR		12		42.821	10.108	73.737	1.00 25.33	
	MOTA		D1 TYR		12		43.338			1.00 23.74	
	MOTA		E1 TYR		12		43.235	8.546		1.00 22.85	
	ATOM		D2 TYR		12		42.183		74.436	1.00 21.93	
	ATOM		E2 TYR		12		42.074	7.793	73.889	1.00 21.99	
	ATOM		Z TYR		12		42.605	7.538	72.640	1.00 22.99	
	ATOM	_	H TYR		12		42.532	6.273	72.109	1.00 18.79	
	MOTA	3097 C			12		42.054	13.567	75.319	1.00 32.74	
	ATOM	3098 0			12		41.986	13.450	76.542	1.00 23.85	
	MOTA	3099 N			13		42.374	14.710	74.720	1.00 26.96	
	ATOM		A GLY		13		42.658	15.900	75.501	1.00 34.92	
	ATOM	3101 C			13		41.452	16.396	76.277	1.00 36.82	
	ATOM	3102 0			13		41.580	17.228	77.176	1.00 34.10	
	MOTA -	3103 N			14		40.279	15.875	75.929	1.00 37.23	
	ATOM	3104 C.			L4		39.031	16.247	76.584	1.00 41.77	
	MOTA	3105 C			L4		37.925	16.406	75.537	1.00 45.82	
	ATOM	3106 C			4		38.110	17.585	74.579	1.00 51.38	
	ATOM	3107 CI			. 4		37.805	18.939	75.241	1.00 57.78	
	ATOM	3108 CI			.4		38.752	19.285	76.388	1.00 58.82	
	MOTA	3109 N2	LYS I	3 1	.4		38.387	20.568	77.070	1.00 55.06	
	ATOM	3110 C	LYS I	3 1	4		38.591	15.226	77.627	1.00 39.50	
	ATOM	3111 o	LYS I	3 1	4		37.546	15.385	78.252	1.00 35.54	
	MOTA	3112 N	TYR I				39.395	14.186	77.815	1.00 40.97	
	ATOM	3113 CA	TYR E	3 1			39.070	13.128	78.768	1.00 40.97	
		3114 CE					38.863	11.827	77.990	1.00 44.15	
•		3115 CG					37.850	11.972	76.876	1.00 44.42	
		3116 CD					38.064	11.372		1.00 42.02	
		3117 CE					37.138		75.634	1.00 41.06	
		3118 CD					36.678	11.530	74.603	1.00 42.76	
		3119 CE					35.748	12.703	77.065	1.00 42.99	
		3120 CZ	TYR B					12.851	76.048	1.00 43.30	
		3121 OH					35.984	12.261	74.816	1.00 45.49	
		3122 C	TYR B				35.066	12.403	73.801	1.00 45.69	
		3123 0	TYR B				40.151	12.944	79.838	1.00 43.48	
		3123 N	ARG B				40.519	11.819	80.167	1.00 41.20	
		3125 CA	ARG B	16			40.647	14.052	80.381	1.00 43.01	
	ATOM :	LA CAL	ם טוא	16	,		41.686	14.012	81.410	1.00 43.70	

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Figure 18-48

ATOM	312	6 C	B ARG	B 16	42.250	15.410	81.663	1.00 49.13
ATOM	312							
ATOM			_		42.656			
					43.858	3 15.624	79.751	. 1.00 55.13
ATOM			E ARG	B 16	44.303	16.549	78.718	1.00 62.87
ATOM	3130	0 C	Z ARG	B 16	44.628			
ATOM	313		H1 ARG					
		_			44.556			
MOTA					45.022	18.600	77.954	1.00 67.15
MOTA	3133	3 C	ARG	B 16	41.093	13.531	82.728	
MOTA	3134	4 0	ARG	B 16	39.882			
ATOM	3135		TYR					
					41.949			
ATOM	3136			B 17	41.494	12.637	84.945	1.00 37.67
MOTA	3137	7 CE	3 TYR	B 17	42.500	11.679	85.584	
ATOM	3138	3 CG	TYR	B 17	42.413			
ATOM	3139							
					42.530			
MOTA	3140				42.502	8.618	83.287	1.00 21.42
ATOM	3141	. CE	2 TYR	B 17	42.258	9.196	85.984	
MOTA	3142	CE	2 TYR	B 17	42.229	7.873	85.556	
ATOM	3143							
					42.355	7.587	84.210	1.00 27.13
MOTA	3144				42.371	6.271	83.796	1.00 19.94
MOTA	3145	С	TYR	B 17	41.377	13.927	85.765	1.00 38.94
MOTA	3146	0	TYR	B 17	41.947	14.951	85.391	
ATOM	3147		PRO					1.00 39.65
					40.647	13.893	86.891	1.00 41.27
MOTA	3148	CD	PRO :	B 18	39.958	12.728	87.462	1.00 43.62
MOTA	3149	CA	PRO 1	B 18	40.448	15.058	87.762	
ATOM	3150	CB	PRO 1	B 18	39.648	14.473	88.928	
MOTA	3151	CG	PRO I					1.00 44.09
				-	40.096	13.015	88.933	1.00 49.22
MOTA	31.52	С	PRO I	-	41.702	15.809	88.221	1.00 45.86
MOTA	3153	0	PRO E	3 18	42.789	15.244	88.317	1.00 45.44
ATOM	3154	N	LYS E	3 19	41.506	17.095	88.507	1.00 48.42
MOTA	3155	CA	LYS E					
					42.535	18.040	88.952	1.00 51.03
ATOM	3156	CB	LYS E		41.973	19.122	89.814	1.00 56.35
MOTA	3157	CG	LYS E	3 19	40.630	18.657	90.563	1.00 65.69
MOTA	3158	CD	LYS E	3 19	40.894	17.441	91.423	
ATOM	3159	CE	LYS E					
					39.602	16.882	91.999	1.00 71.85
MOTA	3160	NZ	LYS E		39.825	15.603	92.731	1.00 72.79
MOTA	3161	С	LYS E	19	43.830	17.593	89.639	1.00 48.62
MOTA	3162	0	LYS E	19	44.912	18.009	89.235	1.00 49.04
ATOM	3163	N	ASN B		43.745			
	3164					16.775	90.678	1.00 43.99
MOTA		CA	ASN B		44.957	16.356	91.375	1.00 43.86
ATOM	3165	CB	ASN B	20	44.740	16.440	92.890	1.00 45.92
MOTA	3166	CG	ASN B	20	44.418	17.848	93.355	1.00 49.44
ATOM	3167	ODI	ASN B		45.194	18.779	93.138	
ATOM	3168	ND2						1.00 47.72
					43.268	18.011	93.999	1.00 49.42
ATOM	3169	С	ASN B	20	45.460	14.960	91.008	1.00 39.49
MOTA	3170	0	ASN B	20	46.496	14.521	91.494	1.00 38.24
MOTA	3171	N	HIS B	21	44.729	14.274	90.140	1.00 36.93
ATOM	3172	CA	HIS B	21				
					45.091	12.923	8 . 723	1.00 33.57
ATOM	3173	CB	HIS B	21	43.948	12.299	87.924	1.00 28.67
ATOM	3174	CG	HIS B	21	44.068	10.817	88.750	1.00 32.14
ATOM	3175	CD2	HIS B	21	44.779	10.076	87.867	
	3176		HIS B					1.00 26.15
ATOM				21	43.431	9.917	89.578	1.00 29.59
MOTA	3177	CEI	HIS B	21	43.743	8.686	89.212	1.00 23.65
ATOM	3178	NE2	HIS B	21	44.560	8.755	88.177	1.00 29.71
ATOM	3179	C	HIS B	21	46.348	12.928	88.852	
								1.00 29.06
ATOM	3180	0	HIS B	21	46.536	13.805	88.015	1.00 24.86
ATCM	3181	N	PRO B	22	47.225	11.937	89.035	1.00 30.50
ATOM	3182	CD	PRO B	22	47.187	10.802	89.976	1.00 31.51
ATOM	3183	CA	PRO B	22	48.446			1 00 20 50
						11.880	88.231	1.00 29.58
ATOM	3184	CB	PRO B	22	49.055	10.549	88.65 6	1.00 33.72
atom	3185	CG	PRO B	22	43.658	10.489	90.124	1.00 31.07
ATOM	3186	С	PRO B	22	48.176	11.950	86.728	1.00 28.52
ATOM	3187	ō	PRO B	22	48.989			
						12.474	85.972	1.00 31.85
atom	3188	N	LEU B	23	47.030	11.435	86.297	1.00 24.47
ATOM	3189		LEU B	23	46.685	11.434	84.874	1.00 27.80
ATOM	3190	CB	LEU B	23	45.933	10.141	84.513	1.00 22.18
atch Atcm	3191		LEU B	23	46.760	8.852	84.556	1.00 29.32
m 1 0.1	J = J =		و نوس		40.700	0.004	04.320	1.00 29.32

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Figure 18-49

MOTA	3192	CD:	L LEU	В	23	45.868	7.628	84.402	1.00	24.94
ATOM	3193	CD2	LEU	В	23	47.805	8.905	83.446	1.00	24.80
	3194	c	LEU		23	45.891	12,638	84.367		27.95
MOTA										
ATOM	3195	၁	LEU	В	23	45.166	12.528	83.373		24.42
MOTA	3196	N	LYS	В	24	46.011	13.793	85.018	1.00	31.01
ATOM	3197	CA	LYS	B	24	45.261	14.946	84.530	1.00	29.40
MOTA	3198	CB	LYS		24	44.934	15.923	85.665		33.03
ATOM	3199	CG	LYS	В	24	45.979	16.969	85.999	1.00	33.54
MOTA	3200	CD	LYS	В	24	47.300	16.397	86.422	1.00	39.10
	3201	CE	LYS		24	48.109	17.466	87.152		45.49
MOTA										
MOTA	3202	NZ	LYS		24	48.224	18.737	86.380		45.95
MOTA	3203	С	LYS	В	24	46.039	15.653	83.425	1.00	30.02
ATOM	3204	0	LYS	В	24	45.508	16.523	82.736	1.00	28.82
	3205		ILE		25		15.262	83.246		25.93
MOTA		N				47.298				
MOTA	3206	CA	ILE		25	48.139	15.858	82.212		29.48
MOTA	3207	CB	ILE	В	25	49.641	15.528	82.409	1.00	33.61
ATOM	3208	CG2	ILE	В	25	50.126	16.033	83.775	1.00	32.27
	3209	CG1			25	49.851	14.014	82.263	1.00	
ATOM										
MOTA	3210.	CD1	. ILE	В	25	51.310	13.584	82.188		36.32
ATOM	3211	С	ILE	B	25	47.784	15.318	80.834	1.00	30.08
ATOM	3212	0	ILE	R	25	47.263	14.210	80.704	1.00	25.37
								79.783		29.19
ATOM	3213	N	PRO		26	48.064	16.101			
ATOM	3214	CD	PRO	В	26	48.650	17.448	79.770		32.47
ATOM	3215	CA	PRO	В	26	47.782	15.673	78.413	1.00	29.52
ATOM	3216	CB	PRO		26	48.103	16.921	77.593	1.00	29.84
							18.046	78.599		36.40
MOTA	3217	CG	PRO		26	47.930				
ATOM	3218	C	PRO	В	26	48.789	14.561	78.137	1.00	27.64
ATOM	3219	0	PRO	В	26	49.920	14.620	78.629	1.00	23.08
ATOM	3220	N	ARG	R	27	48.403	13.557	77.360	1.00	23.09
								77.072		23.00
ATOM	3221	CA	ARG		27	49.326	12.469			
ATOM	3222	CB	ARG	В	27	48.987	11.264	77.962	1.00	26.21
ATOM	3223	CG	ARG	В	27	49.101	11.617	79.449	1.00	17.03
ATOM	3224	CD		В	27	48.663	10.507	80.416	1.00	26.83
								80.502		22.99
ATOM	3225	NE	ARG		27	49.586	9.375			
MOTA	3226	CZ	ARG	В	27	49.444	8.220	79.856		25.06
ATOM	3227	NH1	ARG	В	27	48.408	8.022	79.059	1.00	17.74
ATOM	3228	NH2	ARG	В	27	50.336	7.253	80.027	1.00	23.38
	3229		ARG		27	49.329	12.097	75.595		22.54
MOTA		Ç								
MOTA	3230	O	ARG		27	50.214	12.526	74.852		21.86
ATOM	3231	N	VAL	В	28	48.352	11.318	75.148	1.00	20.64
ATOM	3232	CA	VAL	В	28	48.337	10.954	73.739	1.00	26.57
	3233	CB	VAL		28	47.242	9.917	73.424		30.92
MOTA										
ATOM	3234		VAL		28	47.195	9.645	71.925		27.04
ATOM	3235	CG2	VAL	В	28	47.535	8.616	74.172	1.00	25.45
ATOM	3236	С	VAL	В	28	48.150	12.189	72.866	1.00	28.02
ATOM	3:37	Š	VAL		28	48.780	12.311	71.808		30.88
										24.30
ATOM	3.38	Ŋ	SER		29	47.298	13.112	73.304		
MOTA	3139	CA	SER	В	29	47.082	14.326	72.523	1.00	29.48
MOTA	3240	CB	SER	В	29	45.939	15.169	73.110	1.00	31.72
	3241	OG	SER		29	46.218	15.614	74.424		34.55
MOTA										
MOTA	3242	С	SER		29	48.379	15.125	72.514		30.81
MOTA	3243	O	SER	В	29	48.680	15.820	71.545	1.00	28.85
ATOM	3244	N	LEU	В	30	49.157	15.003	73.589	1.00	29.63
	.3245	CA	LEU		30	50.427	15.721	73.679		31.59
MOTA										
ATOM	3246	CB	LEU		30	51.046	15.593	75.079		29.49
MOTA	3247	CG	LEU	В	30	52.066	16.660	75.513	1.00	34.37
ATOM	3248	CD1	LEU	В	30	52.937	16.083	76.610	1.00	30.15
						52.951	17.098	74.357		32.90
ATOM	3249		LEU		30		_			
ATOM	3250	С	LEU	В	30	51.371	15.085	72.672		25.90
MOTA	3251	2	LEU	В	30	52.052	15.777	71.913	1.00	25.10
	3252	N	LEU		31	51.404	13.756	72.675		22.10
ATOM								71.764		25.52
ATOM	3253	CA	LEU		31	52.268	13.013			
ATOM	3254	CB	LEU	В	31	51.966	11.514	71.842		26.41
ATOM	3255	CG	LEU	В	31	53.066	10.524	71.441	1.00	28.93
	3256		LEU		31	52.425	9.198	71.042	1,00	23.69
ATOM								70.300	1 00	30.41
ATOM	3257	CDZ	LEU	5	31	53.873	11.049	10.300	1.00	20.47

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ATOM 3258 C LEU 3 31	•									
ATOM 3259 O LEU B 31 52.940 13.851 69.614 1.00 21.03 ATOM 3261 CA LEU B 32 50.741 13.481 69.933 1.00 21.27 91 ATOM 3262 CB LEU B 32 48.841 13.798 68.408 1.00 27.91 ATOM 3263 CG LEU B 32 48.841 13.798 68.408 1.00 27.30 ATOM 3264 CD1.LEU B 32 46.699 12.504 68.321 1.00 31.60 ATOM 3265 CD2 LEU B 32 46.699 12.504 68.321 1.00 31.60 ATOM 3267 C LEU B 32 46.699 12.504 68.321 1.00 26.90 ATOM 3267 C LEU B 32 50.835 15.317 68.242 1.00 26.90 ATOM 3267 C LEU B 32 50.835 15.317 68.242 1.00 26.90 ATOM 3267 C LEU B 32 50.835 15.317 68.242 1.00 26.90 ATOM 3269 CA ARG B 33 50.945 16.282 99.111 1.00 28.19 ATOM 3269 CA ARG B 33 50.945 16.282 99.111 1.00 28.19 ATOM 3269 CA ARG B 33 50.945 16.282 99.111 1.00 28.19 ATOM 3271 CG ARG B 33 48.887 18.740 69.904 1.00 41.33 ATOM 3271 CG ARG B 33 48.887 18.740 69.904 1.00 41.33 ATOM 3273 CD ARG B 33 46.977 19.713 70.970 1.00 47.67 ATOM 3275 NH1 ARG B 33 46.977 19.931 70.924 1.00 56.24 ATOM 3275 NH1 ARG B 33 46.977 19.931 70.924 1.00 56.24 ATOM 3275 NH1 ARG B 33 46.977 19.931 70.924 1.00 56.24 ATOM 3277 C ARG B 33 50.926 18.500 69.912 1.00 60.10 ATOM 3277 C ARG B 33 50.926 18.500 69.912 1.00 60.10 ATOM 3277 C ARG B 33 50.926 18.500 69.912 1.00 60.10 ATOM 3279 N PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3278 C PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD1 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD1 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.20 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.30 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.30 ATOM 3283 CD2 PIE B 34 55.096 16.176 70.920 1.00 30.30 ATOM 3283 CD2 PIE B 34 55.096 16.176	MOTA	3258	С	LEU E	3 3 1	52.010	13.489	70.335	1.00	25.38
ATOM 3260 N LEU B 32		3259		LEH F	3 31		13.851	69.614	1.00	21.03
ATOM 3261 CA LEU B 32										
ATOM 3262 CB LEU B 32										
ATOM 3264 COL LEU B 32										
ATOM 3265 CD1 LEU B 32	MOTA									
ATOM 3265 CD2 LEU B 32	ATOM	3263	CG	LEU E	3 32	48.195	12.419	68.614	1.00	27.30
ATOM 3265 CD2 LEU B 32	ATOM	3264	CD1	L_LEU E	3 3 2	46.699	12.504	68.321	1.00	31.60
ATOM 3267 O LEU B 32		3265					11.391	67.708	1.00	26.90
ATOM 3268 N ARG B 33	-						•			
ATOM 3268 N ARG B 33										
ATOM 3269 CA ARG B 33 50.962 17.660 68.865 1.00 31.77 ATOM 3271 CG ARG B 33 50.952 18.601 69.930 1.00 34.22 ATOM 3272 CD ARG B 33 48.887 18.740 69.904 1.00 40.33 ATOM 3273 NC ARG B 33 48.887 18.740 69.904 1.00 47.67 ATOM 3273 NC ARG B 33 46.977 19.931 70.970 1.00 47.67 ATOM 3275 NH1 ARG B 33 46.977 19.931 70.970 1.00 67.04 ATOM 3275 NH1 ARG B 33 46.977 19.931 70.924 1.00 60.10 ATOM 3275 NH1 ARG B 33 46.977 29.929 68.845 1.00 63.11 ATOM 3276 NH2 ARG B 33 45.011 20.652 69.965 1.00 63.11 ATOM 3277 C ARG B 33 55.028 18.580 68.097 1.00 30.12 ATOM 3278 N PHE B 34 53.147 17.012 69.694 1.00 30.70 ATOM 3280 CA PHE B 34 55.096 16.176 70.920 1.00 30.40 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 30.42 ATOM 3282 CB PHE B 34 56.556 16.358 71.248 1.00 28.56 ATOM 3283 CD1 PHE B 34 56.556 16.358 71.248 1.00 28.56 ATOM 3284 CD2 PHE B 34 56.556 16.358 71.248 1.00 28.88 ATOM 3286 CC PHE B 34 58.831 15.530 71.246 1.00 33.71 ATOM 3287 CP PHE B 34 59.265 16.669 71.887 1.00 28.88 ATOM 3288 C PHE B 34 59.265 16.669 71.887 1.00 28.15 ATOM 3288 C PHE B 34 59.265 16.669 71.287 1.00 33.71 ATOM 3289 C PHE B 34 59.265 16.669 71.287 1.00 33.71 ATOM 3280 C PHE B 34 59.265 16.669 71.887 1.00 28.15 ATOM 3280 C PHE B 34 59.265 16.669 71.887 1.00 28.15 ATOM 3280 C PHE B 34 59.265 16.669 71.887 1.00 28.15 ATOM 3280 C PHE B 34 59.265 16.669 71.887 1.00 28.15 ATOM 3290 N LYS B 35 55.202 16.583 68.460 1.00 33.71 ATOM 3290 N LYS B 35 55.202 16.583 68.460 1.00 33.71 ATOM 3290 N LYS B 35 55.202 16.583 68.460 1.00 33.73 ATOM 3291 CA LYS B 35 55.205 16.669 71.887 1.00 48.03 ATOM 3292 CB LYS B 35 55.076 18.929 66.454 1.00 32.97 ATOM 3293 CC LYS B 35 55.076 18.929 66.454 1.00 32.97 ATOM 3299 N ALS B 35 55.076 18.929 66.454 1.00 32.97 ATOM 3300 CA ASP B 36 53.823 16.426 65.560 1.00 33.73 ATOM 3301 CB ASP B 36 53.823 16.426 65.541 1.00 34.33 ATOM 3302 CB ASP B 36 53.670 18.929 66.454 1.00 32.90 ATOM 3303 CD ASP B 36 53.670 18.929 66.454 1.00 32.90 ATOM 3304 OD ASP B 36 54.764 19.099 63.534 1.00 36.44 ATOM 3307 N ALA B 37 55.559 20.655 67.383 1.00 36.44 AT										
ATOM 3271 CG ARG B 33	MOTA	3268	N	ARG E						
ATOM 3271 CB ARG B 33	ATOM	. 3269	CA	ARG E	33	50.962	17.660	68.865	1.00	31.77
ATOM 3271 CG ARG B 33 48.887 18.740 69.904 1.00 40.33 ATOM 3272 CD ARG B 33 48.20 19.713 70.970 1.00 47.67 ATOM 3273 NE ARG B 33 46.977 19.931 70.970 1.00 47.67 ATOM 3273 NE ARG B 33 46.977 19.931 70.970 1.00 47.67 ATOM 3275 NH1 ARG B 33 46.997 20.929 68.845 1.00 63.11 ATOM 3275 NH1 ARG B 33 46.997 20.929 68.845 1.00 63.11 ATOM 3276 NH2 ARG B 33 45.011 20.652 69.965 1.00 63.81 ATOM 3277 C ARG B 33 52.076 17.791 68.852 1.00 30.30 ATOM 3278 O ARG B 33 52.076 17.791 68.852 1.00 30.20 ATOM 3278 O ARG B 33 53.028 18.580 68.097 1.00 30.20 ATOM 3280 CA PHE B 34 53.147 17.012 69.694 1.00 30.20 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 30.40 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 30.40 ATOM 3283 CDI PHE B 34 57.001 17.515 71.885 1.00 26.92 ATOM 3284 CDZ PHE B 34 57.001 17.515 71.885 1.00 26.92 ATOM 3288 CDI PHE B 34 58.346 17.684 72.206 1.00 28.88 ATOM 3285 CEI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3288 CDI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3288 CDI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3288 CDI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3288 CDI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3288 CDI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3289 O PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3289 O PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3289 O PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3299 N LYS B 35 52.20 16.583 68.460 1.00 33.71 ATOM 3290 N LYS B 35 55.202 16.583 68.460 1.00 33.71 ATOM 3290 N LYS B 35 55.202 16.583 68.460 1.00 33.71 ATOM 3290 N LYS B 35 55.204 14.880 66.753 1.00 34.33 ATOM 3291 CA LYS B 35 55.204 14.880 66.753 1.00 34.33 ATOM 3292 CB LYS B 35 55.015 18.82 66.753 1.00 34.33 ATOM 3299 N ASP B 36 53.833 16.426 65.606 1.00 32.75 ATOM 3299 N ASP B 36 53.833 16.426 65.606 1.00 32.75 ATOM 3300 CA ASP B 36 53.833 16.426 65.606 1.00 32.37 ATOM 3300 CA ASP B 36 53.833 16.426 65.606 1.00 32.30 ATOM 3301 CB ASP B 36 53.659 19.901 66.459 1.00 37.66 ATOM 3300 CB ASP B 36 53.659 19.901 66.459 1.00 37.66 ATOM 3300 CB ASP B 36 54.694 19.099 63.534 1.00 36.		3270	CB	ARG E	33	50.395	18.601	69.930	1.00	34.22
ATOM 3272 CD ARG B 33								69.904		
ATOM 3273 NE ARG B 33 46.977 19.931 70.924 1.00 56.24 ATOM 3275 NH1 ARG B 33 46.997 20.929 68.845 1.00 60.10 ATOM 3275 NH1 ARG B 33 46.997 20.929 68.845 1.00 63.11 ATOM 3276 NH2 ARG B 33 45.011 20.652 69.965 1.00 63.81 ATOM 3277 C ARG B 33 52.076 17.791 68.852 1.00 30.12 ATOM 3278 O ARG B 33 52.076 17.791 68.852 1.00 30.12 ATOM 3278 N PHE B 34 53.028 18.580 68.097 1.00 30.20 ATOM 3280 CA PHE B 34 53.147 17.012 69.694 1.00 30.70 ATOM 3281 CB PHE B 34 55.06 16.176 70.920 1.00 30.40 ATOM 3281 CB PHE B 34 56.556 16.358 71.248 1.00 28.56 ATOM 3283 CD1 PHE B 34 56.556 16.358 71.248 1.00 28.56 ATOM 3284 CD2 PHE B 34 56.556 16.358 71.248 1.00 28.85 ATOM 3284 CD2 PHE B 34 58.831 15.537 70.932 1.00 28.88 ATOM 3286 CE2 PHE B 34 58.831 15.537 70.932 1.00 28.88 ATOM 3287 CZ PHE B 34 59.655 16.669 71.887 1.00 28.15 ATOM 3288 C PHE B 34 55.202 16.583 68.460 1.00 33.78 ATOM 3287 CZ PHE B 34 55.202 16.583 68.460 1.00 33.74 ATOM 3287 CZ PHE B 34 55.202 16.583 68.460 1.00 33.74 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 33.74 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 33.74 ATOM 3289 C PHE B 35 55.202 16.583 68.460 1.00 33.74 ATOM 3291 CA LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3292 CB LYS B 35 55.204 14.880 66.753 1.00 34.93 ATOM 3292 CB LYS B 35 55.141 12.423 67.414 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 32.97 ATOM 3294 CD LYS B 35 55.055 15.141 12.423 67.414 1.00 32.97 ATOM 3299 C B LYS B 35 55.055 15.842 67.414 1.00 32.97 ATOM 3293 CG LYS B 35 55.056 16.066 67.704 71.00 41.43 ATOM 3295 CE LYS B 35 55.056 15.066 67.704 71.00 41.43 ATOM 3295 CE LYS B 35 55.056 16.066 67.704 71.00 32.93 ATOM 3297 C LYS B 35 55.056 16.066 67.704 71.00 32.32 ATOM 3290 N ASP B 36 53.468 17.365 64.552 1.00 33.39 ATOM 3290 N ASP B 36 53.468 17.366 67.504 71.00 32.32 ATOM 3300 CA ASP B 36 53.666 53.468 17.366 67.504 1.00 32.32 ATOM 3303 ODI ASP B 36 53.666 53.468 17.366 67.504 1.00 33.480 ATOM 3303 ODI ASP B 36 53.667 18.892 66.569 1.00 33.38 ATOM 3303 ODI ASP B 36										
ATOM 3274 CZ ARC B 33 46.330 20.505 69.912 1.00 60.10 ATOM 3275 NH1 ARG B 33 46.997 20.929 68.845 1.00 63.11 ATOM 3276 NH2 ARG B 33 45.011 20.652 69.965 1.00 63.11 ATOM 3277 C ARG B 33 52.476 17.791 68.852 1.00 30.20 ATOM 3278 O ARG B 33 52.476 17.791 68.852 1.00 30.20 ATOM 3278 O ARG B 34 53.028 NB.580 68.097 1.00 30.20 ATOM 3280 CA PHE B 34 53.147 17.012 69.694 1.00 30.70 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 30.46 ATOM 3282 CG PHE B 34 55.096 16.176 70.920 1.00 30.46 ATOM 3283 CDI PHE B 34 56.556 16.358 71.248 1.00 28.85 ATOM 3283 CDI PHE B 34 57.481 15.373 70.932 1.00 28.85 ATOM 3285 CEI PHE B 34 57.481 15.373 70.932 1.00 28.85 ATOM 3286 CE2 PHE B 34 59.265 16.689 71.246 1.00 31.47 ATOM 3287 CZ PHE B 34 59.265 16.689 71.246 1.00 31.47 ATOM 3280 C PHE B 34 59.265 16.689 71.246 1.00 31.47 ATOM 3280 C PHE B 34 59.265 16.689 71.246 1.00 31.47 ATOM 3280 C PHE B 34 55.006 16.76 70.920 1.00 33.78 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 31.47 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 31.47 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 33.78 ATOM 3290 N LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3291 CA LYS B 35 55.294 14.880 66.753 1.00 34.33 ATOM 3292 CB LYS B 35 55.294 14.880 66.753 1.00 34.33 ATOM 3292 CB LYS B 35 54.580 11.066 67.047 1.00 41.43 ATOM 3295 CB LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CB LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CB LYS B 35 55.151 15.842 65.608 1.00 34.33 ATOM 3296 N Z LYS B 35 55.151 15.842 65.608 1.00 34.33 ATOM 3297 C LYS B 35 55.151 15.842 65.608 1.00 34.33 ATOM 3298 C ASP B 36 53.823 16.426 65.502 1.00 34.33 ATOM 3298 C ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3300 CA ASP B 36 53.651 1.10 66 67.047 1.00 44.04 ATOM 3301 CB ASP B 36 53.651 1.10 66 67.047 1.00 44.04 ATOM 3301 CB ASP B 36 53.651 1.10 66 67.047 1.00 44.04 ATOM 3301 CB ASP B 36 53.651 1.10 66 67.047 1.00 35.78 ATOM 3301 CB ASP B 36 53.651 1.10 8.99 63.534 1.00 32.32 ATOM 3300 CB ASP B 36 53.651 1.10 9.99 63.534 1.00 33.640 ATOM 3301 CB ASP B 36 53.651 1.10 9.99 63.53										
ATOM 3275 NH1 ARC B 33										
ATOM 3276 NH2 ARG B 33	ATOM									
ATOM 3278 C ARG B 33 53.476 17.791 68.852 1.00 30.12 ATOM 3279 N PHE B 34 53.028 18.580 68.097 1.00 30.20 ATOM 3280 CA PHE B 34 54.600 17.060 69.774 1.00 30.70 ATOM 3281 CB PHE B 34 54.600 17.060 69.774 1.00 30.70 ATOM 3282 CG PHE B 34 54.600 17.060 69.774 1.00 29.42 ATOM 3283 CD1 PHE B 34 56.556 16.176 70.920 1.00 30.46 ATOM 3284 CD2 PHE B 34 56.556 16.358 71.248 1.00 28.56 ATOM 3285 CE1 PHE B 34 57.001 17.515 71.885 1.00 28.56 ATOM 3286 CE2 PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3286 CE2 PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3287 CZ PHE B 34 59.265 16.689 71.887 1.00 28.15 ATOM 3288 C PHE B 34 59.265 16.689 71.887 1.00 28.15 ATOM 3289 O PHE B 34 55.202 16.583 68.460 1.00 33.78 ATOM 3290 N LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3291 CA LYS B 35 55.294 14.80 66.753 1.00 34.33 ATOM 3292 CB LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3294 CD LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CE LYS B 35 55.05 11.004 67.205 1.00 41.43 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 33.33 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 36.33 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 36.33 ATOM 3299 N ASP B 36 53.468 17.882 65.608 1.00 33.33 ATOM 3299 C A SP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3300 CA ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3301 CB ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3302 CB ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3303 CD ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3301 CB ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3301 CB ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3301 CB ALA B 37 55.599 20.659 67.383 1.00 38.64 ATOM 3301 CB ALA B 37 55.599 19.901 65.429 1.00 37.66 ATOM 3311 C ALA B 37 55.599 19.901 65.429 1.00 37.66 ATOM 3312 N MET B 38 59.341 17.488 67.474 1.00 37.62 ATOM 3313 CB MET B 38 59.341 17.488 67.474 1.00 37.62 ATOM 3314 CB MET B 38 59.341 17.488 67.474 1.00 37.62 ATOM 3313 C MET B 38 59.341 17.486 67.404 1.00 38.67 ATOM 3311 C MET B 38 59.341 17.486 67.404 1.00 38.67 ATOM	ATOM	3275	NHI	L ARG B	33	46.997	20.929	68.845	1.00	63.11
ATOM 3278	ATOM	3276	NH2	ARG B	33	45.011	20.652	69.965	1.00	63.81
ATOM 3279 N PHE B 34 53.147 17.012 69.694 1.00 30.20 ATOM 3280 CA PHE B 34 55.096 16.176 70.920 1.00 30.46 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 30.46 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 28.56 ATOM 3283 CD1 PHE B 34 55.096 16.358 71.248 1.00 28.56 ATOM 3283 CD1 PHE B 34 57.001 17.515 71.885 1.00 26.92 ATOM 3285 CE1 PHE B 34 57.001 17.515 71.885 1.00 26.92 ATOM 3285 CE1 PHE B 34 58.831 15.530 71.246 1.00 28.56 ATOM 3286 CE2 PHE B 34 58.836 17.684 72.206 1.00 28.156 ATOM 3287 CZ PHE B 34 58.831 15.530 71.246 1.00 31.47 ATOM 3287 CZ PHE B 34 59.265 16.689 71.887 1.00 28.155 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 33.78 ATOM 3290 N LYS B 35 54.580 11.066 67.047 1.00 33.78 ATOM 3291 CA LYS B 35 54.580 11.066 67.047 1.00 34.93 ATOM 3292 CB LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CE LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CE LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3297 C LYS B 35 55.3070 11.004 67.205 1.00 44.04 ATOM 3299 N ASP B 36 53.823 11.984 66.345 1.00 44.04 ATOM 3299 N ASP B 36 53.823 11.984 66.345 1.00 35.37 ATOM 3299 C LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3301 CB ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3301 CB ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3300 CA ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3300 CA ASP B 36 53.845 1.00 64.698 1.00 43.03 ATOM 3301 CB ASP B 36 53.845 1.00 64.698 1.00 43.03 ATOM 3301 CB ASP B 36 53.845 1.00 64.698 1.00 43.03 ATOM 3300 CA ASP B 36 53.845 1.00 64.698 1.00 35.78 ATOM 3300 CA ASP B 36 53.845 1.00 64.698 1.00 36.845 ATOM 3301 CB ASP B 36 53.845 1.00 64.698 1.00 36.845 ATOM 3301 CB ASP B 36 53.845 1.00 66.656 66.013 1.00 36.845 ATOM 3310 C ALA B 37 55.559 20.659 67.383 1.00 33.85 ATOM 3310 C ALA B 37 55.559 20.659 67.938 1.00 36.845 A		3277	С	ARG B	33	52.476	17.791	68.852	1.00	30.12
ATOM 3279 N PHE B 34 53.147 17.012 69.694 1.00 30.70 ATOM 3280 CA PHE B 34 54.6600 17.060 69.774 1.00 29.42 ATOM 3281 CB PHE B 34 55.096 16.176 70.920 1.00 30.46 ATOM 3281 CD PHE B 34 56.556 16.358 71.248 1.00 28.56 ATOM 3283 CD1 PHE B 34 57.001 17.515 71.885 1.00 26.92 ATOM 3284 CD2 PHE B 34 57.001 17.515 71.885 1.00 26.92 ATOM 3285 CE1 PHE B 34 57.481 15.373 70.932 1.00 28.88 ATOM 3285 CE2 PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3286 CE2 PHE B 34 59.265 16.689 71.287 1.00 31.47 ATOM 3287 CZ PHE B 34 59.265 16.689 71.887 1.00 33.78 ATOM 3289 C PHE B 34 55.202 16.583 68.466 1.00 33.78 ATOM 3289 C PHE B 34 55.202 16.583 68.466 1.00 33.78 ATOM 3291 CA LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3291 CA LYS B 35 55.294 14.880 66.753 1.00 34.93 ATOM 3292 CB LYS B 35 55.141 12.423 67.414 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 32.97 ATOM 3295 CE LYS B 35 55.3070 11.004 67.205 1.00 44.04 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 33.73 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 42.56 ATOM 3300 CA ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.66 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.42 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.66 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.66 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.68 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.68 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.68 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.68 ATOM 3301 CB ASP B 36 54.764 19.099 63.534 1.00 37.68 ATOM 3310 CB ASP B 38 60.841 18.391 67.784 1.00									1.00	30.20
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ATOM 3284 CD2 PHE B 34 57.481 15.373 70.932 1.00 28.88 ATOM 3285 CE1 PHE B 34 58.881 15.530 71.246 1.00 31.47 ATOM 3286 CE2 PHE B 34 58.881 15.530 71.246 1.00 31.47 ATOM 3287 CZ PHE B 34 59.265 16.689 71.887 1.00 28.15 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 33.71 ATOM 3290 N LYS B 35 54.770 15.413 67.999 1.00 33.71 ATOM 3291 CA LYS B 35 54.770 15.413 67.999 1.00 38.63 ATOM 3292 CB LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3293 CG LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3294 CD LYS B 35 54.580 11.066 67.047 1.00 41.43 ATOM 3295 CE LYS B 35 54.580 11.066 67.047 1.00 41.43 ATOM 3297 C LYS B 35 52.335 11.984 66.345 1.00 60.09 ATOM 3298 O LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 33.39 ATOM 3299 N ASP B 36 53.468 17.365 64.552 1.00 33.39 ATOM 3300 CA ASP B 36 53.468 17.365 64.552 1.00 33.39 ATOM 3301 CB ASP B 36 51.812 18.544 62.461 1.00 42.56 ATOM 3303 OD ASP B 36 51.812 18.544 62.461 1.00 43.04 ATOM 3303 OD ASP B 36 51.812 18.544 62.461 1.00 42.56 ATOM 3304 OD2 ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 55.594 19.901 65.429 1.00 36.21 ATOM 3309 CB ALA B 37 55.595 19.901 65.429 1.00 36.25 ATOM 3311 O ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3312 N MET B 38 59.158 17.059 66.013 1.00 36.25 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 36.25 ATOM 3314 CB MET B 38 59.341 17.438 67.474 1.00 37.66 ATOM 3315 C MET B 38 59.341 17.438 67.474 1.00 37.66 ATOM 3316 C MET B 38 59.341 17.438 67.474 1.00 37.66 ATOM 3317 C MET B 38 59.341 17.438 67.474 1.00 37.66 ATOM 3310 C MET B 38 59.559 17.262 63.130 1.00 35.86 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 36.25 ATOM 3312 N MET B 38 59.559 17.262 63.130 1.00 35.86 ATOM 3313 C MET B 38 59.559 17.262 63.130 1.00 35.86 ATOM 3310 C MET B 38 59.559 17.262 63.130 1.00 35.86 ATOM 3310 C MET B 38 59.559 17.262 63.130 1.00 35.86 ATOM 3310 C MET B 38 59.559 17.262 63.130 1.00 35.86	MOTA	3283	CD1	PHE B	34	57.001	17.515	71.885	1.00	26.92
ATOM 3285 CEI PHE B 34 58.346 17.684 72.206 1.00 28.15 ATOM 3286 CE2 PHE B 34 58.831 15.530 71.246 1.00 31.47 ATOM 3287 CZ PHE B 34 59.265 16.689 71.887 1.00 32.87 ATOM 3288 C PHE B 34 55.202 16.583 68.460 1.00 33.78 ATOM 3289 O PHE B 34 56.049 17.259 67.873 1.00 33.71 ATOM 3290 N LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3291 CA LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3292 CB LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3293 CG LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3294 CD LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3295 CE LYS B 35 54.580 11.066 67.047 1.00 41.43 ATOM 3295 CE LYS B 35 53.070 11.004 67.205 1.00 44.09 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3298 O LYS B 35 55.869 16.061 64.752 1.00 33.39 ATOM 3299 N ASP B 36 53.488 17.365 64.552 1.00 33.39 ATOM 3300 CA ASP B 36 53.488 17.365 64.552 1.00 32.32 ATOM 3301 CB ASP B 36 53.488 17.365 64.552 1.00 32.32 ATOM 3302 CG ASP B 36 51.812 18.544 62.461 1.00 29.17 ATOM 3304 OD2 ASP B 36 51.812 18.544 62.461 1.00 29.17 ATOM 3305 C ASP B 36 51.812 18.544 62.461 1.00 32.32 ATOM 3307 N ALA B 37 54.694 19.099 63.534 1.00 42.56 ATOM 3309 CB ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3301 C ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 37.42 ATOM 3312 N MET B 38 59.158 17.250 66.3130 1.00 37.42 ATOM 3313 CA NET B 38 59.158 17.250 66.3130 1.00 36.44 ATOM 3315 CG MET B 38 59.158 17.250 66.3130 1.00 37.68 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 37.68 ATOM 3312 N MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3313 CA NET B 38 59.158 17.262 63.130 1.00 35.86 ATOM 3316 C MET B 38 59.541 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 59.541 17.426 63.035 1.00 35.86 ATOM 3310 N ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 37.68 ATOM 3312 C MET B 38 59.541 17.262 63.130 1.00 35.86 ATOM 3313 C MET B 38 59.659 17.262 63.130 1.00 35.86 ATOM 3310 C ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3322 C B ASN B 39 57.470 17.742 63.035 1.00 46.207			CD2	PHE B	34	57.481	15.373	70.932	1.00	28.88
ATOM 3286 CE2 PHE B 34 58.831 15.530 71.246 1.00 31.47 ATOM 3287 CZ PHE B 34 59.265 16.689 71.887 1.00 28.15 ATOM 3288 C PHE B 34 59.265 16.689 71.887 1.00 28.15 ATOM 3289 C PHE B 34 55.202 16.583 68.460 1.00 33.78 ATOM 3290 N LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3291 CA LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3292 CB LYS B 35 54.770 15.413 67.999 1.00 28.65 ATOM 3293 CG LYS B 35 55.294 14.880 66.753 1.00 34.33 ATOM 3292 CB LYS B 35 55.294 14.880 66.753 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CE LYS B 35 54.580 11.006 67.047 1.00 41.43 ATOM 3295 CE LYS B 35 55.015 11.004 67.205 1.00 44.04 ATOM 3295 CE LYS B 35 55.015 15.842 65.608 1.00 60.09 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 33.39 ATOM 3300 CA ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 52.015 17.800 64.658 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3303 OD1 ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3303 OD1 ASP B 36 54.371 18.590 64.578 1.00 42.56 ATOM 3306 CA ASP B 36 54.371 18.590 64.578 1.00 30.480 ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 30.864 ATOM 3301 C ALA B 37 55.554 20.226 65.924 1.00 30.56 ATOM 3310 C ALA B 37 55.554 20.226 65.924 1.00 30.56 ATOM 3311 C ALA B 37 55.559 20.659 67.383 1.00 34.80 ATOM 3311 C ALA B 37 55.559 20.659 67.383 1.00 37.68 ATOM 3311 C ALA B 37 55.559 20.659 67.383 1.00 37.68 ATOM 3311 C ALA B 37 55.559 20.659 67.383 1.00 37.68 ATOM 3311 C ALA B 37 55.559 20.659 67.383 1.00 37.68 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3311 C ALA B 37 55.559 20.659 67.383 1.00 37.68 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.582 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.582 ATOM 3311 C ALA B 37 57.675 20.776 64.950 1.00 30.582 ATOM 33								72 206		
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ATOM 3291 CA LYS B 35 55.294 14.880 66.753 1.00 34.33 ATOM 3292 CB LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3293 CG LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3294 CD LYS B 35 55.141 12.423 67.414 1.00 44.93 ATOM 3295 CE LYS B 35 54.580 11.066 67.047 1.00 41.43 ATOM 3295 CE LYS B 35 53.070 11.004 67.205 1.00 44.04 ATOM 3296 NZ LYS B 35 52.335 11.984 66.345 1.00 60.09 ATOM 3299 N ASP B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3301 CB ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3302 CG ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3303 OD1 ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3305 C ASP B 36 54.371 18.859 64.578 1.00 6.14 ATOM 3306 O ASP B 36 54.371 18.859 64.578 1.00 6.14 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3309 CB ALA B 37 55.554 20.226 67.383 1.00 34.84 ATOM 3310 C ALA B 37 55.554 20.226 67.383 1.00 38.54 ATOM 3311 O ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3312 N MET B 38 59.158 17.059 66.013 1.00 36.25 ATOM 3312 N MET B 38 59.158 17.059 66.013 1.00 37.62 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 37.62 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 37.62 ATOM 3316 CD MET B 38 59.158 17.059 66.013 1.00 37.62 ATOM 3317 CE MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3317 CE MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3318 C MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3318 C MET B 38 59.659 17.228 67.300 1.00 30.56 ATOM 3319 O MET B 38 59.659 17.228 67.300 1.00 30.58 ATOM 3319 O MET B 38 59.659 17.228 67.300 1.00 35.82 ATOM 3320 N ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 33	ATOM	3289	0	PHE B	34	56.049	17.259			
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ATOM 3292 CB LYS B 35 54.684 13.509 66.454 1.00 32.97 ATOM 3293 CG LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3294 CD LYS B 35 55.141 12.423 67.414 1.00 34.93 ATOM 3295 CE LYS B 35 54.580 11.006 67.205 1.00 44.04 ATOM 3296 NZ LYS B 35 53.070 11.004 67.205 1.00 44.04 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3298 O LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.824 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3304 OD2 ASP B 36 51.812 18.544 66.461 1.00 79.17 ATOM 3305 C ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3306 O ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3308 CA ALA B 37 54.694 19.091 63.534 1.00 32.40 ATOM 3309 CB ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 37.66 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3316 CB MET B 38 59.158 17.059 66.013 1.00 36.24 ATOM 3317 CE MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3318 C MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3319 O MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3319 O MET B 38 59.659 17.262 63.035 1.00 35.82 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 35.82 ATOM 3320 N ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 47.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 47.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 47.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 47.75 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 47.75 ATOM 332		3291	CA	LYS B	35	55.294	14.880	66.753	1.00	34.33
ATOM 3293 CG LYS B 35								66.454	1.00	32.97
ATOM 3294 CD LYS B 35 54.580 11.066 67.047 1.00 41.43 ATOM 3295 CE LYS B 35 53.070 11.004 67.205 1.00 44.04 ATOM 3296 NZ LYS B 35 52.335 11.984 66.345 1.00 60.09 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3298 O LYS B 35 55.869 16.061 64.752 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3303 OD1 ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3304 OD2 ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3305 C ASP B 36 51.111 19.897 64.043 1.00 44.34 ATOM 3306 O ASP B 36 54.371 18.590 64.578 1.00 66.14 ATOM 3307 N ALA B 37 54.694 19.096 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 54.694 19.096 65.777 1.00 34.80 ATOM 3300 CB ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 37.66 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 37.42 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 37.42 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3316 CD MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3317 CE MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3318 C MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 CD MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3317 CE MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3318 C MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 CD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 59.659 17.262 63.130 1.00 32.89 ATOM 3318 C MET B 38 59.659 17.262 63.130 1.00 32.89 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 32.89 ATOM 3321 CA ASN B 39 57.470 17.742 63.035 1.00 32.89 ATOM 3321 CA ASN B 39 57.470 17.742 63.035 1.00 42.75 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 42.75 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 42.75 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.75 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 46.20										
ATOM 3295 CE LYS B 35 53.070 11.004 67.205 1.00 44.04 ATOM 3296 NZ LYS B 35 52.335 11.984 66.345 1.00 60.09 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3298 N ASP B 36 55.869 16.061 64.752 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.822 63.661 1.00 42.56 ATOM 3303 OD1 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3304 OD2 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3305 C ASP B 36 51.111 19.897 64.043 1.00 4.34 ATOM 3305 C ASP B 36 54.371 18.590 64.578 1.00 61.14 ATOM 3306 O ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.559 20.659 67.383 1.00 38.54 ATOM 3310 C ALA B 37 55.559 20.659 67.383 1.00 38.54 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 37.66 ATOM 3312 N MET B 38 59.554 18.635 65.541 1.00 37.42 ATOM 3312 N MET B 38 59.341 17.438 67.474 1.00 36.25 ATOM 3314 CB MET B 38 59.341 17.438 67.474 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.42 ATOM 3316 CD MET B 38 59.341 17.438 67.784 1.00 38.07 ATOM 3317 CE MET B 38 59.341 17.438 67.474 1.00 38.07 ATOM 3318 C MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3318 C MET B 38 59.659 17.228 67.300 1.00 30.56 ATOM 3318 C MET B 38 59.659 17.228 67.300 1.00 30.98 ATOM 3318 C MET B 38 59.659 17.228 67.300 1.00 30.98 ATOM 3319 O MET B 38 59.659 17.228 67.300 1.00 35.82 ATOM 3321 CA ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322										
ATOM 3296 NZ LYS B 35 52.335 11.984 66.345 1.00 60.09 ATOM 3297 C LYS B 35 55.015 15.842 65.608 1.00 35.78 ATOM 3298 O LYS B 35 55.015 15.842 65.608 1.00 33.39 ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3304 OD2 ASP B 36 51.111 19.897 64.043 1.00 4.34 ATOM 3305 C ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3306 O ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3309 CB ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.559 20.666 67.383 1.00 38.54 ATOM 3311 O ALA B 37 55.599 20.626 65.924 1.00 37.66 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 36.25 ATOM 3314 CB MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3318 C MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3319 O MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3311 CA ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57										
ATOM 3297 C LYS B 35	ATOM									
ATOM 3298 O LYS B 35	ATOM	3296	NZ							
ATOM 3298 O LYS B 35	ATOM	3297	С	LYS B	35	55.015	15.842	65.608	1.00	35.78
ATOM 3299 N ASP B 36 53.823 16.426 65.602 1.00 32.32 ATOM 3300 CA ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3304 OD2 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3305 C ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3306 O ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3307 N ALA B 37 54.694 19.099 63.534 1.00 32.40 ATOM 3308 CA ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3309 CB ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.554 20.226 65.924 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 37.68 ATOM 3319 O MET B 38 59.659 17.226 63.130 1.00 30.98 ATOM 3319 O MET B 38 59.659 17.226 63.130 1.00 30.98 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3321 CA ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.757 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.757 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.0		3298	0	LYS B	35	55.869	16.061	64.752	1.00	33.39
ATOM 3300 CA ASP B 36 53.468 17.365 64.552 1.00 36.31 ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.812 18.544 62.461 1.00 79.17 ATOM 3304 OD2 ASP B 36 51.11 19.897 64.043 1.00 4.34 ATOM 3305 C ASP B 36 54.764 19.099 63.534 1.00 4.34 ATOM 3306 O ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3309 CB ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3311 O ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 59.158 17.059 66.013 1.00 36.25 ATOM 3314 CB MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 30.956 ATOM 3318 C MET B 38 62.093 17.226 67.300 1.00 30.98 ATOM 3318 C MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3318 C MET B 38 59.659 17.262 63.130 1.00 30.98 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 30.98 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3321 CA ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57			N	ASP B	36	53.823	16.426	65.602	1.00	32.32
ATOM 3301 CB ASP B 36 52.015 17.800 64.698 1.00 42.56 ATOM 3302 CG ASP B 36 51.617 18.822 63.661 1.00 43.03 ATOM 3303 OD1 ASP B 36 51.812 18.544 62.461 1.00 '9.17 ATOM 3304 OD2 ASP B 36 51.111 19.897 64.043 1.00 4.34 ATOM 3305 C ASP B 36 54.371 18.590 64.578 1.00 '6.14 ATOM 3306 O ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3311 O ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET 3 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 37.68 ATOM 3318 C MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3318 C MET B 38 58.639 17.226 63.130 1.00 30.98 ATOM 3319 O MET B 38 58.639 17.226 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57 ATOM 3322 CB ASN B 39 57.321 17.262 61.661 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.742 63.035 1.00 47.57 ATOM 3322 CB ASN B 39 57.470 17.684 ATOM 3322 CB ASN B 39									1.00	36.31
ATOM 3302 CG ASP B 36										
ATOM 3303 OD1 ASP B 36										
ATOM 3304 OD2 ASP B 36 51.111 19.897 64.043 1.00 4.34 ATOM 3305 C ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3306 O ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3310 C ALA B 37 55.554 20.226 65.924 1.00 37.66 ATOM 3311 O ALA B 37 56.959 20.659 67.383 1.00 38.67 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3314 CB MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 CB MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET B 38 62.093 17.226 67.300 1.00 30.98 ATOM 3318 C MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57	ATOM									
ATOM 3305 C ASP B 36 54.371 18.590 64.578 1.00 6.14 ATOM 3306 O ASP B 36 54.764 19.099 63.534 1.00 32.40 ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3309 CB ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3310 C ALA B 37 56.959 19.01 65.429 1.00 37.66 ATOM 3311 O ALA B 37 56.959 19.01 65.429 1.00 37.66 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3314 CB MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET B 38 62.093 17.226 67.300 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57	MOTA									
ATOM 3306 O ASP B 36	ATOM	3304	OD2	ASP B	36					
ATOM 3306 O ASP B 36	MOTA	3305	С	ASP B	36	54.371	18.590	64.578	1.00	`6.14
ATOM 3307 N ALA B 37 54.694 19.061 65.777 1.00 34.80 ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3309 CB ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3310 C ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3316 SD MET B 38 60.841 18.391 67.784 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 37.68 ATOM 3318 C MET B 38 60.841 18.391 67.784 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.222 67.300 1.00 30.98 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.86 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57				ASP B	36	54.764	19.099	63.534	1.00	32.40
ATOM 3308 CA ALA B 37 55.554 20.226 65.924 1.00 36.82 ATOM 3309 CB ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3310 C ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.158 17.059 66.013 1.00 37.68 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 37.68 ATOM 3318 C MET B 38 62.093 17.228 67.300 1.00 30.98 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 35.86 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57		_						65.777	1.00	34.80
ATOM 3310 C ALA B 37 55.599 20.659 67.383 1.00 38.54 ATOM 3310 C ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3317 CE MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3318 C MET B 38 62.093 17.228 67.300 1.00 30.98 ATOM 3319 O MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57										
ATOM 3310 C ALA B 37 56.959 19.901 65.429 1.00 37.66 ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET B 38 62.093 17.223 67.300 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.581 1.00 47.57										
ATOM 3311 O ALA B 37 57.675 20.776 64.950 1.00 30.56 ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET B 38 62.093 17.223 67.300 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.588 1.00 47.57	ATOM									
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ATOM 3312 N MET B 38 57.346 18.635 65.541 1.00 37.42 ATOM 3313 CA MET B 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET B 38 62.093 17.223 67.300 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57	ATOM	3311	0	ALA 3	37	57.675	20.776	64.950		
ATOM 3313 CA MET 3 38 58.670 18.192 65.107 1.00 36.25 ATOM 3314 CB MET 3 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET 3 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET 3 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET 3 38 62.093 17.226 67.300 1.00 30.98 ATOM 3318 C MET 3 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET 8 38 58.639 17.690 63.663 1.00 35.86 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 32.69 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57		3312	N	MET B	38	57.346	18.635	65.541		
ATOM 3314 CB MET B 38 59.158 17.059 66.013 1.00 36.44 ATOM 3315 CG MET B 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET B 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET B 38 62.093 17.228 67.300 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57							18,192	65.107	1.00	36.25
ATOM 3315 CG MET 3 38 59.341 17.438 67.474 1.00 37.68 ATOM 3316 SD MET 3 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET 3 38 62.093 17.228 67.300 1.00 30.98 ATOM 3318 C MET 3 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET 8 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57									1.00	36 44
ATOM 3316 SD MET 3 38 60.841 18.391 67.784 1.00 38.07 ATOM 3317 CE MET 3 38 62.093 17.223 67.300 1.00 30.98 ATOM 3318 C MET 3 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET 8 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57										
ATOM 3317 CE MET 3 38 62.093 17.223 67.300 1.00 30.98 ATOM 3318 C MET 5 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET 5 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN 5 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN 5 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN 5 39 58.156 18.108 60.688 1.00 47.57										
ATOM 3317 CE MET B 38 62.093 17.223 67.300 1.00 30.98 ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.20	MOTA									
ATOM 3318 C MET B 38 58.639 17.690 63.663 1.00 35.86 ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.757	ATOM	3317	CE .	MET 3						
ATOM 3319 O MET B 38 59.659 17.262 63.130 1.00 32.69 ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 47.57			С	MET B	38	58.639	17.690	63.663	1.00	35.86
ATOM 3320 N ASN B 39 57.470 17.742 63.035 1.00 35.82 ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.20						59.659	17.262	63.130		
ATOM 3321 CA ASN B 39 57.321 17.262 61.661 1.00 42.75 ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.20										
ATOM 3322 CB ASN B 39 58.156 18.108 60.688 1.00 46.20										
ATOM 2222 GC 200 P 30 57 570 10 543 60 591 1.00 47 57										
2262 AC 201 B 20 67 670 10 642 BD 591 1.00 47 57	MOTA									
	MOTA	3323	CG	ASN B	39	57.670	19.543	00.591	1.00	4/.5/

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ATON	4 3324 OD1 ASN	B 39	56.524	19.801	60.212	1.00 48.78
ATON			58.540			
ATOM						
ATOM		_	57.759			
			58.465		60.639	1.00 35.75
ATOM			57.332	14.997	62.535	1.00 34.64
ATOM	F 3329 CA LEU	B 40	57.700	13.590		
ATOM	I 3330 CB LEU	B 40	58.347			
ATOM			59.595			
ATOM			60.148		-	
ATOM						
			60.646			
MOTA			56.549		62.264	1.00 37.58
ATOM			56.637	11.438	62.573	
ATOM	3336 N ILE	B 41	55.476	13.131	61.663	
MOTA			54.340	12.290	61.314	
MOTA			53.445	11.991		
ATOM					62.536	1.00 35.21
ATOM			52.793	13.271	63.047	1.00 31.89
			52.367	10.980	62.141	1.00 32.68
ATOM			51.470	10.550	63.285	1.00 36.46
ATOM	3342 C ILE	B 41	53.492	12.937	60.229	1.00 37.52
MOTA	3343 O ILE	B 41	53.352	14.157	60.183	1.00 40.24
ATOM	3344 N ASP	B 42	52.943	12.114	59.345	1.00 39.55
ATOM	3345 CA ASP		52.094	12.615		1.00 39.33
ATOM	3346 CB ASP				58.273	1.00 45.30
			52.569	12.119	56.901	1.00 45.93
ATOM			53.972	12.584	56.564	1.00 47.09
ATOM	3348 OD1 ASP		54.244	13.799	56.686	1.00 46.60
ATOM	3349 OD2 ASP	B 42	54.797	11.736	56.162	1.00 45.16
MOTA	3350 C ASP	B 42	50.677	12.134	58.524	1.00 45.15
ATOM	3351 O ASP	B 42	50.467	11.051	59.069	1.00 47.06
ATOM	3352 N GLU		49.707	12.944	58.121	
ATOM	3353 CA GLU		48.303		50.121	1.00 48.13
ATOM	3354 CB GLU			12.618	58.312	1.00 50.50
			47.441	13.637	57.571	1.00 53.54
ATOM			45.961	13.505	57.840	1.00 59.52
ATOM	3356 CD GLU I		45.155	14.518	57.065	1.00 64.03
ATOM	3357 OE1 GLU E	3 43 .	43.914	14.535	57.215	1.00 68.54
ATOM	3358 OE2 GLU E	3 43	45.765	15.298	56.301	1.00 66.95
ATOM	3359 C GLU E	3 43	47.972	11.205	57.836	1.00 47.81
ATOM	3360 O GLU E		47.092	10.547	58.390	1.00 49.67
ATOM	3361 N LYS E		48.690	10.744	56.817	
ATOM	3362 CA LYS E	-				1.00 46.21
MOTA	3363 CB LYS E	_	48.484	9.409	56.251	1.00 48.28
			49.207	9.311	54.894	1.00 49.96
MOTA	3364 CG LYS B		49.639	7.903	54.470	1.00 52.18
MOTA	3365 CD LYS B		50.970	7.532	55.127	1.00 61.03
MOTA	3366 CE LYS B	44	51.399	6.095	54.844	1.00 62.80
ATOM	3367 NZ LYS B	44	50.511	5.098	55.510	1.00 65.34
ATOM	3368 C LYS B	44	48.899	8.249	57.161	1.00 45.92
ATOM	3369 O LYS B		48.418	7.127	57.009	1.00 41.30
ATOM	3370 N GLU B					
ATOM	3371 CA GLU B	-	49.797	8.517	581.100	1.00 42.18
			50.268	7.486	59.014	1.00 38.41
ATOM			51.684	7.812	59.468	1.00 33.73
ATOM	3373 CG GLU B	45	52.694	7.887	58.351	1.00 37.58
MOTA	3374 CD GLU B	45	53.998	8.504	58.813	1.00 34.34
MOTA	3375 OE1 GLU B	45	53.997	9.699	59.176	1.00 38.04
ATOM	3376 OE2 GLU B	45	55.020	7.799	58.821	1.00 33.37
ATOM	3377 C GLU B	45	49.368	7.403	60.238	1.00 36.86
ATOM	3378 O GLU B	45			61 033	
ATOM			49.461	6.462	61.032	1.00 34.98
		46	48.489	8.386	60.386	1.00 30.86
MOTA	3380 CA LEU B	46 .	47.608	8.438	61.545	1.00 30.65
MOTA	3381 CB LEU B	46	47.501	9.889	62.019	1.00 32.74
ATOM	3382 CG LEU B	46	46.642	10.163	63.250	1.00 34.76
ATOM	3383 CD1 LEU B	46	47.189	9.379	64.425	1.00 32.24
MOTA	3384 CD2 LEU B	46	46.639	11.656	63.548	1.00 32.24
ATOM	3385 C LEU B	46	46.212			
ATOM	3386 O LEU B	46		7.861	61.318	1.00 31.36
			45.530	8.218	60.363	1.00 31.78
atom	3387 N ILE B	47	45.801	6.957	62.203	1.00 31.18
ATCM	3388 CA ILE B	47	44.479	6.338	62.139	1.00 29.36
ATOM	3389 CB ILE B	47	44.564	4.802	62.258	1.00 28.62
					-	- -

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MOTA	3390) c	32 ILE	B 47		43.161	4.205	62.407	1.00 28.80
ATOM	3391		31 ILE						
						45.266		61.028	
MOTA	3392		Ol ILE			45.419	2.722	61.054	
MOTA	3393	C	ILE :	B 47		43.659	6.875	63.303	1.00 32.22
MOTA	3394	0	ILE I		•	44.063		64.461	
							_		
MOTA	3395		LYS			42.514		62.999	
MOTA	3396	C.A	LYS I	3 48		41.662	8.037	64.040	1.00 32.37
ATOM	3397	CE	LYS I	3 48		40.517	8.840	63.414	
MOTA	3398					39.607	9.514		
								64.430	
ATOM	3399					38.535	10.361	63.747	1.00 44.38
ATOM	3400	CE	LYS E	3 48	-	37.65 7	11.074	64.768	1.00 45.91
ATOM	3401	N2	LYS E	3 48		38.451	11.991	65.643	
ATOM	3402		LYS E						
						41.095	6.943	64.937	
MOTA	3403		LYS E			40.524	5.962	64.457	1.00 26.24
MOTA	3404	N	SER E	3 4,9		41.260	7.121	66.244	1.00 27.89
ATOM	3405	CA	SER E	49		40.770	6.168	67.232	1.00 25.17
ATOM	3406	СВ							
						41.146	6.639	68.642	1.00 24.41
MOTA	3407	OG	SER E	49		42.539	6.858	68.777	1.00 31.79
MOTA	3408	С	SER B	49		39.248	6.054	67.160	1.00 29.07
MOTA	3409	0	SER B	49.		38.565	7.034	66.879	1.00 28.47
ATOM	3410	Ň	ARG B						
						38.723	4.859	67.409	1.00 26.13
ATOM	3411	CA	ARG B	50		37.278	4.658	67.430	1.00 24.24
MOTA	3412	CB	ARG B	50		36.810	3.700	66.323	1.00 25.03
MOTA	3413	CG	ARG B			37.231	2.233	66.507	1.00 26.54
								– .	
MOTA	3414	CD	ARG B			36.570	1.340	65.452	1.00 26.21
MOTA	3415	NE	ARG B	50		37.006	-0.058	65.504	1.00 25.13
ATOM	3416	CŻ	ARG B	50		36.700	-0.924	66.468	1.00 26.09
ATOM	3417		l ARG B	50		35.941	-0.558	67.497	1.00 23.42
		NH2							
MOTA	3418			50		37.157	-2.168	66.402	1.00 23.91
ATOM	3419	С	ARG B	50		36.937	4.037	68.775	1.00 23.83
MOTA	3420	0	ARG B	50		37.782	3.392	69.403	1.00 21.60
MOTA	3421	N	PRO B	51		. 35.700	4.223	69.243	1.00 22.99
MOTA	3422	CD	PRO B	51					
						34.554	4.962	68.688	1.00 25.09
MOTA	3423	ÇA	PRO B	51		35.338	3.628	70.530	1.00 25.48
MOTA	3424	CB	PRO B	51		33.949	4.217	70,802	1.00 26.32
ATOM	3425	CG	PRO B	51		33.936	5.503	69.953	1.00 28.65
ATOM	3426	C	PRO B	51		35.264	2.118	70.325	
									1.00 26.73
ATOM	3427	0	PRO B	51		35.142	1.646	69.194	1.00 18.87
ATOM	3428	N	ALA B	52		35.355	1.359	71.408	1.00 23.64
MOTA	3429	CA	ALA B	52		35.237	-0.083	71,291	1.00 23.27
ATOM	3430	CB	ALA B	52		35.811	-0.757	72.521	1.00 26.31
	3431	č							
ATOM			ALA B	52		33.733	-0.324	71.223	1.00 25.25
ATOM	3432	0	ALA B	52		32.950	0.515	71.677	1.00 22.78
ATOM	3433	N	THR B	53		33.321	-1.447	70.651	1.00 22.77
ATOM	3434	CA	THR B	53		31.900	-1.760	70.596	1.00 26.90
	3435	CB	THR B	53					
MOTA						31.567	-2.732	69.456	1.00 30.00
MOTA	3436	OG1	THR B	53		32.305	-3.950	69.642	1.00 25.59
MOTA	3437	CG2	THR B	53		31.917	-2.117	68.103	1.00 23.33
MOTA	3438	С	THR B	53		31.579	-2.445	71.916	1.00 30.41
MOTA	3439	ō	THR B	53				72.609	
						32.484	-2.917		1.00 26.13
MOTA	3440	N	LYS B	54		30.300	-2.504	72.268	1.00 29.12
MOTA	3441	ÇA	LYS B	54		29.909	-3.140	73.514	1.00 30.24
MOTA	3442	CB	LYS B	54		28.396	-3.027	73.720	1.00 32.78
	3443	CG	LYS B					75.131	
MOTA				54		27.947	-3.351		1.00 34.85
MOTA	3444	CD	LYS B	54		26.445	-3.204	75.268	1.00 41.13
MOTA	3445	CE	LYS B	54		26.008	-3.366	76.709	1.00 43.39
MOTA	3446	NZ	LYS B	54		26.464	-2.257	77.582	1.00 47.39
ATOM	3447	c	LYS B	54		30.329	-4.603	73.442	1.00 29.10
MOTA	3448	0	LYS B	54		30.779	-5.183	74.430	1.00 26.71
MOTA	3449	N	GLU B	55		30.196	-5.187	72.256	1.00 23.97
MOTA	3450	CA	GLU B	55		30.577	-6.577	72.032	1.00 28.08
		CB	GLU B						
ATCM	3451			55		30.288	-6.965	70.579	1.00 24.82
MOT	3452	CG	GLU B	55		30.671	-8.400	70.237	1.00 33.40
TOM	3453	CD	GLU B	55		30.453	-8.737	68.767	1.00 38.49
TOM	3454	OE1	GLU B	55		30.638	-9.913	68.394	1.00 41.24
	3455		GLU B	55		30.101		67.984	1.00 40.02
MOT	ر ر په ر		OHO B	J J		JU. 101	-7.833	-1.304	1.00 40.02

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Figure 18-53

```
32.066
                                                               1.00 25.82
         3456
               C
                    GLU B
                           55
                                             -6.808
                                                      72.345
 ATOM
         3457
                           55
 ATOM
                    GLU B
                                    32.429
                                             -7.765
                                                      73.033
                                                               1.00 23.83
 MOTA
         3458
               N
                    GLU B
                           56
                                    32.931
                                             -5.938
                                                      71.835
                                                               1.00 25.04
                           56
                                    34.365
         3459
                    GLU B
ATOM
               CA
                                             -6.079
                                                      72.093
                                                               1.00 25.30
ATOM
         3460
               CB
                    GLU B
                           56
                                    35.141
                                             -5.003
                                                      71.334
                                                               1.00 24.31
 ATOM-
         3461
                CG
                    GLU B
                           56
                                    34.866
                                             -5.039
                                                      69.836
                                                               1.00 32.15
                    GLU B
MOTA
         3462
               CD
                           56
                                    35.512
                                             -3.903
                                                      69.073
                                                               1.00 31.43
         3463
               OE1
                    GLU B
                           56
                                    35.486
                                             -2.759
                                                      69.568
                                                               1.00 28.54
ATOM
                                    36.012
                                                      67.959
MOTA
         3464
               OE2
                    GLU B
                           56
                                             -4.147
                                                               1.00 28.89
                                             -5.988
                                                      73.595
 MOTA
         3465
               C
                    .GLU B
                           56
                                    34.653
                                                               1.00 28.88
ATOM
         3466
               0
                    GLU B
                           56
                                    35.450
                                             -6.766
                                                      74.137
                                                               1.00 25.07
         3467
               N
                    LEU B.
                           57
                                    33.996
                                             -5.050
                                                      74.272
                                                               1.00 24.52
ATOM
                                                      75.702-
                    LEU B
                           57
                                    34.203
MOTA
         3468
               CA
                                             -4.891
                                                               1.00 27.34
         3469
               CB
                    LEU B
                           57
                                    33.416
                                             -3.694
                                                      76.231
                                                               1.00 22.79
ATOM
                    LEU B
                           57
                                    33.859
                                                      75.722
         3470
               CG
                                             -2.320
                                                               1.00 23.57
MOTA
               CD1
                           57
                                                      76.366
ATOM
         3471
                   LEU B
                                    33.008
                                             -1.247
                                                               1.00 22.27
ATOM
         3472
               CD2
                   LEU B
                           57
                                    35.342
                                             -2.089
                                                      76.061
                                                               1.00 17.24
         3473
               С
                    LEU B
                           57
                                    33.785
                                             -6.144
                                                      76.452
                                                               1.00 26.92
MOTA
        3474
                           57
                                             -6.568
                                                      77.396
                                                               1.00 24.06
               O
                   LEU B
                                    34.458
ATOM
                                                      76.029
                                                               1.00 23.35
MOTA
        3475
               Ν
                    LEU B
                           58
                                    32.670
                                             -6.732
        3476
                   LEU B
                           58
                                    32.154
                                             -7.931
                                                      76.674
                                                               1.00 25.60
MOTA
               CA
                                    30.718
        3477
                   LEU B
                           58
                                             -8.207
                                                      76.221
                                                               1.00 28.50
               CB
ATOM
                   LEU B
                                    29.734
                                                      76.649
                                                               1.00 30.91
ATOM
        3478
               CG
                           58
                                             -7.110
               CD1
                   LEU B
                           58
                                    28.323
                                             -7.468
                                                      76.212
                                                               1.00 28.93
ATOM
        3479
        3480
               CD2
                   LEU B
                           58
                                    29.794
                                             -6.945
                                                      78.157
                                                               1.00 33.44
ATOM
                                    33.027
                                            -9.153
        3481
                   LEU B
                                                      76.446
                                                               1.00 24.59
                           58
MOTA
               C
        3482
               0
                   LEU B
                           58
                                    32.760
                                            -10.216
                                                      76.991
                                                               1.00 19.76
ATOM
                   LEU B
                           59
                                    34.065
                                            -9.006
                                                      75.630
                                                               1.00 23.99
MOTA
        3483
               N
                                                               1.00 25.11
        3484
               CA
                   LEU B
                           59
                                    34.988 -10.108
                                                      75.411
MOTA
                                                               1.00 21.64
                                            -9.757
                                                      74.332
MOTA
        3485
               CB
                   LEU B
                           59
                                    36.018
                   LEU B
                           59
                                    35.483
                                             -9.652
                                                      72.905
                                                               1.00 24.24
ATOM
        3486
               CG
                           59
                                    36.585
                                                      71.975
        3487
               CD1
                   LEU B
                                            -9.177
                                                               1.00 24.25
MOTA
                           59
                                                      72.468
                                                               1.00 19.91
        3488
               CD2 LEU B
                                    34.957 ~11.014
MOTA
                           59
                                    35.699 -10.371
                                                               1.00 23.84
        3489
               С
                   LEU B
                                                      76.733
ATOM
MOTA
        3490
                   LEU B
                           59
                                    36.150 -11.489
                                                      76.992
                                                               1.00 19.39
        3491
                   PHE B
                           60
                                    35.793
                                            -9.344
                                                      77.577
                                                               1.00 21.80
               N
ATOM
                                    36.462
                                                               1.00 23.08
                                            -9.510
                                                      78.876
                   PHE B
                           60
ATOM
        3492
               CA
                                            -8.770
ATOM
        3493
               CB
                   PHE B
                           60
                                    37.809
                                                      78.908
                                                               1.00 18.22
        3494
               CG
                   PHE B
                           60
                                    38.544
                                            -8.906
                                                      80.230
                                                               1.00 21.72
MOTA
        3495
                   PHE B
                           60
                                    38.975 -10.157
                                                      80.680
                                                               1.00 19.23
               CD1
ATOM
                                    38.757
                                            -7.791
                                                      81.048
                                                               1.00 17.75
ATOM
        3496
               CD2
                   PHE B
                           60
        3497
               CE1
                   PHE B
                           60
                                    39.602 -10.301
                                                      81.927
                                                               1.00 18.80
MOTA
                                                      82.297
                                                               1.00 19.23
        3498
              CE2
                   PHE B
                          60
                                    39.384
                                            -7.923
ATOM
                                                      82.737
                                                               1.00 16.10
                          60
                                    39.807
                                            -9.184
ATOM
        3499
              CZ
                   PHE B
        3500
                   PHE B
                           60
                                    35.648
                                            -9.069
                                                      80.083
                                                               1.00 21.58
MOTA
                                   35.508
                                                      81.040
                                                               1.00'22.21
        3501
              0
                   PHE B
                          60
                                            -9.822
MOTA
                                            -7.847
                                                      80.055
                                                               1.00 20.65
ATOM
        3502
              N
                   HIS B
                          61
                                   35.128
        3503
                   HIS B
                          61
                                   34.362
                                            -7.336
                                                      81.184
                                                               1.00 23.32
MOTA
              CA
        3504
              СВ
                   HIS B
                          61
                                   34.422
                                            -5.807
                                                      81.229
                                                               1.00 27.60
ATOM
                                                      81.440
                                                               1.00 31.83
                                            -5.259
ATOM
        3505
              CG
                  HIS B
                          61
                                   35.800
                                                              1.00 26.86
        3506
              CD2 HIS B
                          61
                                   36.466
                                            -4.940
                                                      82.575
ATOM
                                   36.669
                                            -5.003
                                                      80.401
                                                               1.00 34.35
        3507
              ND1
                  HIS B
                          61
MOTA
                                   37.810
                                            -4.546
                                                      80.887
                                                               1.00 34.78
ATOM
        3508
              CE1
                  HIS B
                          61
                                                              1.00 36.27
        3509
              NE2 HIS B
                          61
                                   37.713
                                            -4.499
                                                      82,204
ATOM
                          61
                                   32.902
                                            -7.775
                                                      81.198
                                                              1.00 28.04
        3510
              С
                   HIS B
ATOM
                                                      80.176
                   HIS B
                          61
                                   32.349
                                            -8.167
                                                              1.00 25.70
       3511
ATCM
              0
                                                              1.00 25.25
                                   32.276
                                            -7.691
                                                      82.367
ATOM
       3512
              N
                   THR B
                          62
                          62
                                   30.882
                                            -8.084
                                                      82.506
                                                              1.00 25.35
ATOM
       3513
              CA
                   THR B
                                                      83.932
                                   30.578
                                                              1.00 25.47
                          62
                                            -8.549
ATOM
       3514
              CB
                  THR B
                                            -7.462
                                                              1.00 28.62
                          62
                                   30.783
                                                      84.843
       3515
              OG1
                  THR B
ATOM
                          62
                                   31.482
                                            -9.701
                                                      84.315
                                                              1.00 21.32
       3516
              CG2
                  THR B
ATOM
                                                              1.00 26.06
                  THR B
                          62
                                   29.931
                                            -6.942
                                                      82.162
ATOM
       3517
              C
                                                              1.00 24.14
                          62
                                   30.287
                                            -5.771
                                                     82.254
       3518
              0
                  THR B
ATOM
       3519
              N
                  GLU B
                          63
                                   28.718
                                            -7.305
                                                     81.759
                                                              1.00 28.01
ATCM
                                                      81.389
                          63
                                   27.681
                                            -6.349
                                                              1.00 30.77
              CA
                  GLU B
       3520
ATCM
                                                     81.114
                                                              1.00 33.97
                          63
                                   26.374
              CB
                                            -7.094
       3521
                  GLU B
ATCM
```

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				•					
ATOM	3522	CG .GLU	B 63		25.213	-6.21	0 00 66	7 7 0	
ATOM	3523	CD GLU			25.189			_	0 41.12
ATOM		OE1 GLU							0 44.4
					24.361		7 78.68		0 42.64
ATOM		OE2 GLU			25.992	-6.64	78.46	5 1.0	0 45.96
ATOM		C GLU			27.436	-5.32	82.49		0 27.29
ATOM	3527	O GLU	B 63		27.381				0 25.13
ATOM	3528	N ASP			27.272			1.0	0 23.13
ATOM		CA ASP							0 24.38
ATOM					27.010				0 29.27
					26.887		86.112	2 1.00	0 36.30
ATOM			B 64		28.022	-6.939	86.198	3 1.00	50.77
MOTA		OD1 ASP	B 64		29.128				52.71
MOTA	3533	OD2 ASP	B 64		27.812				
ATOM	3534		B 64		28.075				51.98
MOTA	3535	O ASP							26.49
	3536				27.768		85.422		18.33
ATOM		N TYR			29.332	-4.373	85.052	1.00	22.75
ATOM	3537	CA TYR I			30.420	-3.435	85.251		19.32
MOTA	3538	CB TYR I	B 65		31.751	-4.186			16.59
ATOM	3539	CG TYR I	B 65		32.949	-3.285			19.19
MOTA	3540	CD1 TYR F			33.033	-2.328			1 19.19
ATOM	3541	CE1 TYR E							21.35
	3542				34.135	-1.489			18.32
ATOM	•	CD2 TYR E			34.004	-3.382	84.456	1.00	18.65
ATOM	3543	CE2 TYR E	_		35.116	-2.544	84.554		21.01
MOTA		CZ TYR E	65		35.172	-1.601	85.573		20.61
ATOM	3545	OH TYR E	65		36.262	-0.775	85.682		
MOTA	3546	C TYR E			30.392				17.77
ATOM	3547	O TYR B				-2.373	84.146		22.01
MOTA	3548				30.399	-1.167	84.421	1.00	18.20
					30.330	-2.815	82.894	1.00	19.49
ATOM	3549	CA ILE B			30.305	-1.870	81.786	1.00	19.68
ATOM	3550	CB ILE B			30.208	-2.592	80.432		23.31
ATOM		CG2 ILE B	66		30.200	-1.571	79.303		21.30
MOTA	3552	CG1 ILE B	66		31.400	-3.541	80.260		27.67
MOTA	3553	CD1 ILE B			32.758	-2.839			
ATOM		C ILE B	66		29.128		80.291		29.29
MOTA		_				-0.909	81.940	1.00	26.99
		_	66		29.294	0.309	81.848	1.00	23.36
MOTA		n asn b	67	- 2	27.939	-1.447	82.198	1.00	24.98
MOTA		CA ASN B	67	- 2	6.782	-0.580	82.363		27.70
ATOM	3558	CB ASN B	67	2	5.492	-1.389	82.580		25.58
MOTA	3559	CG ASN B	67		5.081	-2.183	81.341		26.91
MOTA	3560	OD1 ASN B	67		5.199	-1.701			
ATOM		ND2 ASN B	67				80.220	1.00	31.48
MOTA		ASN B			4.572	-3.387	81.545		23.80
			67		6.982	0.401	83.513	1.00	25.34
MOTA) ASN B	67	2	6.524	1.539	83.448	1.00	22.53
ATOM	3564 N	1 THR B	68	2	7.664	-0.031	84.568		23.65
ATOM	3565 C	CA THR B	68	2	7.903	0.863	85.696		25.25
MOTA	3566 0	B THR B	68		8.516	0.119	86.891		
MOTA		G1 THR B	68						29.08
ATOM		G2 THR B			7.561	-0.826	87.396		25.94
			68		8.894	1.100	88.002	1.00	22.90
ATOM	3569 C		68		8.818	2.009	85.287	1.00	25.91
MOTA	3570 O		68	2	8.576	3.156	85.661	1.00	28.47
MOTA	3571 N	LEU B	69	2	9.861	1.702	84.519	1.00	
MOTA	3572 C	A LEU B	69		0.788	2.729	84.054	1.00	
MOTA	3573 °C		69		1.915	2.122			
MOTA	3574 C		69				83.201	1.00	
					2.960	1.231	83.889	1.00	
ATOM			69		1.006	0.786	82.859	1.00	22.57
MOTA		D2 LEU B	69	3:	3.643	2.000	85.008	1.00	23.20
MOTA	3577 C	LEU B	69		0.036	3.764	83.229	1.00	
MOTA	3578 O	LEU B	69).190	4.966	83.444		
ATOM	3579 N		70					1.00	
					3.218	3.290	82.294	1.00	
TOM	3580 CZ		70		.449	4.181	81.434	1.00 2	25.87
MOTA	3581 CI		70	27	.660	3.371	80.401	1.00 2	24.80
MOTA	3582 CC		70	28	.531	2.511	79.490	1.00 3	
MOT	3583 SI	MET B	70		.592	1.599	78.227	1.00	
TOM	3584 CE		70		.922	2.986	77.245		
TOM	3585 C	MET B	70					1.00	
					.489	5.062	82.242	1.00 2	
MOT		MET B	70		.391	6.273	82.009	1.00 2	4.09
MOT	3587 N	GLU B	71	26	.786	4.458	83.194	1.00 2	8.21
									-

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30036	3588	CA	GLU	ъ	71	25.837	5.207	84.008	1.00 27.45
MOTA									
MOTA	3589	CB	GLU	В	71	25.014	4.268	84.889	1.00 30.70
ATOM	3590	CG	GLU	В	71	24.072	5.005	85.832	1.00 33.61
	3591	CD	GLU		71	23.044	5.867	85.096	1.00 37.51
MOTA									
ATOM	3592	OE1	GLU	В	71	22.333	6.638	85.773	1.00 35.47
MOTA	3593	OE2	GLU	В	71	22.934	5.769	83.849	1.00 31.03
	3594	¢	GLU		71	26.559	6.209	84.887	1.00 29.15
MOTA									
MOTA	3595	0	GLU	В	71	26.115	7.341	85.035	1.00 23.96
MOTA	3596	N	ALA	В.	72	27.671	5.781	85.481	1.00 27.76
		CA	ALA		72	28.454	6.662	86.340	1.00 27.58
MOTA	3597								
MOTA	3598	CB	ALA	В	72	29.663	5.920	86.909	1.00 23.24
ATOM	3599	С	ALA	В	72	28.924	7.886	85.563	1.00 28.07
	3600	0	ALA		72	28.895	8.999	86.079	1.00 23.22
MOTA	_								1.00 26.40
ATOM	3601	N	GLU		73	29.356	7.684	84.322	
MOTA	3602	CA	GLU	В	73	29.846	8.801	83.529	1.00 29.06
MOTA	3603	CB	GLU	В	73	30.658	8.314	82.325	1.00 29.48
					73	31.162	9.466	81.443	1.00 31.00
MOTA	3604	CG	GLU						
MOTA	3605	CD	GLU	В	73	31.938	9.009	80.216	1.00 34.37
MOTA	3606	OE1	GLU	В	73	33.059	8.461	80.356	1.00 28.41
	3607		GLU		73	31.419	9.203	79.100	1.00 30.59
ATOM									
MOTA	3608	С	GLU		73	28.744	9.734	83.045	1.00 31.92
ATOM	3609	0	GLU	В	73	28.894	10.951	83.104	1.00 35.69
	3610	N	ARG	R	74	27.633	9.186	82.570	1.00 33.53
MOTA			ARG		74	26.583	10.067	82.081	1.00 38.64
ATOM	3611	CA							
MOTA	3612	CB	ARG	В	74	25.456	9.280	81.403	1.00 39.90
MOTA	3613	CG	ARG	В	74	24.448	8.706	82.363	1.00 46.67
	3614	CD	ARG	R	74	23.174	8.311	81.646	1.00 47.53
MOTA					74	22.076	8.153	82.594	1.00 55.58
MOTA	3615	NE	ARG						
MOTA	3616	CZ	ARG	В	74	21.609	9.136	83.362	1.00 56.04
MOTA	3617	NH1	ARG	В	74	22.142	10.351	83.297	1.00 58.93
	3618		ARG		74	20.601	8.910	84.192	1.00 53.62
MOTA					74	26.008	10.914	83.222	1.00 35.84
ATOM	3619	Ç	ARG						
ATOM	3620	0	ARG	В	74	25.778	12.107	83.048	1.00 29.44
ATOM	3621	N	SER	В	75	25.794	10.302	84.386	1.00 31.02
	3622	CA	SER		75	25.243	11.014	85.539	1.00 31.99
MOTA								86.510	1.00 34.47
ATOM	3623	CB	SER		75	24.592	10.038		
ATOM	3624	OG	SER	В	75	25.581	9.228	87.123	1.00 34.33
ATOM	3625	С	SER	В	75	26.339	11.754	86.288	1.00 35.42
		ō	SER		75	26.060	12.555	87.180	1.00 33.45
ATOM	3626								1.00 33.25
ATOM	3627	И	GLN		76	27.584	11.473	85.922	
ATOM	3628	CA	GLN	В	76	28.739	12.082	86.565	1.00 35.61
	3629	CB	GLN	В	76	28.818	13.572	86.241	1.00 30.11
ATOM		CG	GLN		76	30.216	14.112	86.390	1.00 39.13
ATOM	3630								
ATOM	3631	CD	GLN	В	76	31.124	13.681	85.248	
ATOM	3632	OE1	GLN	В	76	31.052	12.546	84.761	1.00 29.21
	3633	NE2	GLN	В	76	31.995	14 583	84.827	1.00 40.93
MOTA					76	28.624	11 892	88.079	1.00 37.88
ATOM	3634	С	GLN						
MOTA	3635	0	GLN	В	76 .	28.901	12 308	88.858	1.00 32.74
ATOM	3636	N	SER	В	77	28.209	10.697	88.488	1.00 34.72
		CA	SER		77	28.047	10.382	89.901	1.00 37.07
MOTA	3637								1.00 39.61
ATOM	3638	CB	SER		77	26.635	10.738	90.371	
ATOM	3639	OG	SER	В	77	25.678	9.941	89.688	1.00 39.03
	3640	С	SER	В	77	28.265	8.897	90.112	1.00 35.95
MOTA						28.177	8.108	89.173	1.00 36.60
ATOM	3641	0	SER		77				1.00 33.03
MOTA	3642	N	VAL	В	78	28.528	8.518	91.355	
	3643	CA	VAL	В	78	28.753	7.124	91.685	1.00 33.41
ATOM		CB	VAL		78	29.742	6.979	92.848	1.00 36.91
MOTE	3644								1.00 34.37
MOTA	3645	CG1	VAL		78	29.955	5.499	93.163	
ATOM	3646	CG2	VAL	В	78	31.055	7.658	92.496	1.00 34.19
	3647	С	VAL		78	27.461	6.431	92.082	1.00 34.93
ATOM			VAL		78	26.897	6.703	93.143	1.00 28.25
MOTA	3648	ე 							1.00 36.73
ATOM	3649	Я	PRO		79	26.971	5.521	91.228	
ATOM	3650	CĐ	PRO	В	79	27.532	5.114	89.930	1.00 37.44
	3651	CA	PRO		79	25.738	4.779	91.493	1.00 38.33
ATOM			PRO		79	25.668	3.826	90.301	1.00 38,68
MOTA	3652	CB	PKU	2				89.201	1.00 37.41
2 TOM	3653	CG	PRO	ರ	79	26.293	4.664	07.201	1.00 37.41

ATOM 3653 CG PRO B 79 26.293 4.664

SUBSTITUTE SHEET (RULE 26)

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MOTA	3654	ı c	PRO	В	79	25.788	4.046	92.834	1.00 36.92
ATOM	3655	5 0	PRO	В	79	26.854	3.648	93.298	1.00 33.03
ATOM	3656	N	LYS		80	24.623	3.881	93.448	1.00 38.43
ATOM	3657	CA	LYS	В	80	24.482	3.206	94.736	1.00 39.73
ATOM	3658				80	23.003	2.871	94.967	1.00 43.33
ATOM	3659				80	22.679	2:129	96.262	1.00 44.60
MOTA	3660				80	21.198	1.742	96.287	1.00 48.09
ATOM	3661		-		80	20.805	1.014	97.559	1.00 50.12
MOTA	3662				80	20.932	1.890	98.760	1.00 53.16
ATOM	3663		LYS		80	25.315	1.928	94.854	1.00 40.35
ATOM	3664		LYS		80	. 25.181	1.011	94.047	1.00 36.67
MOTA	3665	N	GLY	В	81	26.173	1.880	95.869	1.00 38.26
ATOM	3666	CA	GLY	В	81	26.996	0.709	96.104	1.00 34.69
MOTA	3667	С	GLY	В	81	28.066	0.407	95.071	1.00 34.63
MOTA	3668	0	GLY	В	81	28.861	-0.513	95.255	1.00 33.92
MOTA	3669	N	ALA	В	82	28.100	1.178	93.992-	1.00 31.26
MOTA	3670	CA	ALA	В	82	. 29.082	0.963	92.936	1.00 34.88
ATOM	3671	CB	ALA	В	82	28.755	1.848	91.751	1.00 23.13
MOTA	3672	С	ALA	В	82	30.517	1.223	93.405	1.00 36.85
ATOM	3673	О	ALA	В	82	31.461	0.580	92.945	1.00 32.17
MOTA	3674	N	ARG	В	83	30.677	2.168	94.323	1.00 36.52
MOTA	3675	CA	ARG	В	83	31.994	2.522	94.830	1.00 38.75
MOTA	3676	CB	ARG	В	83	31.865	3.616	95.885	1.00 40.24
MOTA	3677	CG	ARG		83	33.187	4.180	96.330	1.00 49.12
MOTA	3678	CD	ARG		83	33.015	5.239	97.404	1.00 53.26
MOTA	3679	NE	ARG		83	34.240	6.010	97.624	1.00 59.30
ATOM	3680	CZ	ARG		83	35.437	5.486	97.883	1.00 61.56
ATOM	3681	NH1			83	35.598	4.170	97.958	1.00, 63.53
ATOM	3682	NH2			83	36.479	6.285	98.073	1.00 62.02
ATOM -	3683	C	ARG		83	32.719	1.326	95.426	1.00 37.75
ATOM	3684	0	ARG		83	33.893	1.094	95.146	1.00 37.18
ATOM	3685 3686	N CA	GLU :		84	32.011 32.581	0.564	96.249	1.00 35.29
MOTA	3687	CB	GLU		84 84	31.876	-0.609 -0.855	96.898 98.236	1.00 35.29
MOTA MOTA	3688	CG	GLU :		84	30.443	-0.383	98.240	1.00 40.14
ATOM	3689	CD	GLU I		84	30.356	1.132	98.293	1.00 46.30
MOTA	3690	OE1			84	29.339	1.690	97.834	1.00 43.07
ATOM	3691	OE2	GLU I		84	31.306	1.762	98.814	1.00 50.07
MOTA	3692	С	GLU I		84	32.527	-1.880	96.055	1.00 32.90
MOTA	3693	0	GLU 1		84	33.371	-2.765	96.193	1.00 28.68
ATOM	3694	N	LYS I	В	85	31.533	-1.984	95.187	1.00 27.12
ATOM	3695	CA	LYS F	В	85	31.412	-3.177	94.361	1.00 30.46
ATOM	3696	CB	LYS E	3	85	29.950	-3.401	93.967	1.00 30.01
MOTA	3697	CG	LYS E	3	85	29.717	-4.643	93.117	1.00 28.40
ATOM	3698	CD	LYS E	3	85	28.234	-4.807	92.775	1.00 32.87
MC A	3699	CE	LYS E		85	28.000	-6.048	91.928	1.00 34.15
A. OM	3700	NZ	LYS E		85	26.582	-6:186	91.507	1.00 35.34
MCIA	3701	C	LYS E		85	32.267	-3.096	93.101	1.00 28.98
ATOM	3702	0	LYS E		85	32.817	-4.098	92.652	1.00 24.69
ATOM	3703	N	TYR E		86	32.391	-1.896	92.550	1.00 27.81
ATOM	3704	CA	TYR B		86	33.141	-1.692		1.00 27.56
ATOM	3705	CB	TYR B		86	32.206	-1.050	90.288	1.00 28.88
MOTA	3706 3707	CG CD1	TYR B		86 86	31.008	-1.927 -3.137	89.951	1.00 31.29
MOTA	3708	CEI	TYR B		86 86	31.178 30.095		89.276	1.00 26.99
MOTA	3709		TYR B		86	29.713	-3.955 -1.553	88.965 90.315	1.00 26.97
MOTA	3710		TYR B		86	28.611	-2.370	90.008	1.00 28.38 · 1.00 24.19
MOTA	3711	CZ	TYR B		86	28.815	-3.569	89.331	1.00 28.46
MOTA	3712		TYR B		86	27.747	-4.379	89.008	1.00 22.70
MOTA MOTA	3713		TYR B		86	34.422	-0.870	91.489	1.00 24.64
	3714		TYR B		86	35.160	-0.645	90.530	1.00 27.19
MOTA MOTA	3715		ASN B		37	34.674	-0.418	92.711	1.00 27.19
ATOM	3716		ASN B		37	35.881	0.341	93.032	1.00 29.30
ATOM ATOM	3717		ASN B		37	37.105	-0.561	92.866	1.00 28.92
ATOM	3718		ASN B		37	38.343	0.019	93.506	1.00 34.72
ATOM	3719		ASN B		37	38.309	0.452	94.659	1.00 38.41
						_			

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3 mOM	3720	ND2	ASN B	87	39.449	0.012	92.775	1.00 3	35.86
ATOM							92.223	1.00 2	
MOTA	3721	С	asn b	87	36.070	1.622			
MOTA	3722	0	ASN B	87	37.194	1.998	91.876	1.00 2	
MOTA	3723	N	ILE B	88	34.956	2.282	91.932	1.00 2	29,43
ATOM	3724	CA	ILE B	88	34.945	3.536	91.196	1.00	30.64
					33.959	3.464	90.027	1.00	
ATOM-	3725	CB	ILE B	88					
ATOM	3726	CG2	ILE B	88	33.821	4.829	89.379	1.00	
MOTA	3727	CG1	ILE B	88	34.421	2.433	89.008	1.00	35.43
	3728	CD1		88	35.684	2.821	88.324	1.00 4	41.80
ATOM						4.669	92.118	1.00	
ATOM	3729	С	ILE B	88	34.483				
MOTA	3730	0	ILE B	88	33.681	4.445	93.024	1.00	
MOTA	3731	N	GLY B.	89	34.977	5.881	91.875	1.00	30.36
	3732	CA	GLY B	89	34.574	7.022	92.686	1.00 2	29.54
ATOM					35.601	7.524	93.685	1.00	
MOTA	3733	С	GLY B	89				1.00	
ATOM	3734	0	GLY B	89	35.497	8.652	94.177		
ATOM	3735	N	GLY B	90	36.583	6.687	94.005	1.00	30.97
ATOM	3736	CA	GLY B	90	37.612	7.086	94.949	1.00	31.03
			GLY B	90	38.655	7.936	94.247	1.00	34 78
ATOM	3737	C						1.00	
ATOM	3738	0	GLY B	90	38.455	8.344	93.103		
ATOM	3739	N	TYR B	91	39.772	8.201	94.915	1.00	
ATOM	3740	CA	TYR B	91	40.820	9.023	94.322	1.00	28.15
		СВ	TYR B	91	41.810	9.463	95.405	1.00 2	27.29
MOTA	3741						96.007	1.00	
MOTA	3742	CG	TYR B	91	42.609	8.330			
ATOM	3743	CD1	TYR B	91	43.738	7.823	95.359	1.00	
MOTA	3744	CE1	TYR B	91	44.456	6.762	95.896	1.00	28.75
	3745	CD2	TYR B	91	42.219	7.741	97.208	1.00	28.35
MOTA				91	42.927	6.680	97.751	1.00	27.58
MOTA	3746	CE2	TYR B					1.00	
ATOM	3747	CZ	TYR B	91	44.043	6.196	97.094		
ATOM	3748	oh	TYR B	91	44.753	5.154	97.637	1.00	
ATOM	3749	С	TYR B	91	41.563	8.271	93.226	1.00	29.27
	3750	o	TYR B	91	42.109	8.874	92,308	1.00	25.22
ATOM					41.568	6.948	93.318	1.00	
MOTA	3751	N	GLU B	92				1.00	
ATOM	3752	CA	GLU B	92	42.286	6.124.	92.350		
ATOM	3753	CB	GLU B	92	42.474	4.726	92.924	1.00	
ATOM	3754	CG	GLU B	92	43.502	3.884	92.221	1.00	29.80
	3755	CD	GLU B	92	43.585	2.500	92.826	1.00	35.34
MOTA						1.645	92.477	1.00	
ATOM	3756	OE1		92	42.742				
MOTA	3757	OE2	GLU B	92	44.475	2.278	93.678	1.00	
ATOM	3758	С	GLU B	92	41.594	6.024	90.997	1.00	23.42
	3759	0	GLU B	92	42.204	6.260	89.962	1.00	20.47
ATOM			ASN B	93	40.314	5.677	91.017	1.00	18.85
MOTA	3760	N					89.795	1.00	
MOTA	3761	CA	ASN B	93	39.534	5.509			
ATOM	3762	CB	ASN B	93	39.165	4.033	89.664	1.00	
ATOM	3763	CG	ASN B	93	40.351	3.120	89.943	1.00	24.78
	3764	OD1		93	41.362	3.160	89.239	1.00	22.35
MOTA			ASN B	93	40.240	2.311	90.987	1.00	13.35
MOTA	3765	ND2					89.944	1.00	
MOTA	3766	С	ASN B	93	38.285	6.362			
MOTA	3767	0	asn b	93	37.183	5.843	90.121	1.00	
ATOM	3768	N	PRO B	94	38.449	7.693	89.887	1.00	26.37
	3769	CD	PRO B	94	39.738	8.389	89.716	1.00	19.35
MOTA				94	37.373	8.676	90.024	1.00	24.59
MOTA	3770	CA	PRO B						
MOTA	3771	CB	PRO B	94	38.147	9.972	90.200		25.95
MOTA	3772	CG	PRO B	94	39.297	9.740	89.223	1.00	
MOTA	3773	C	PRO B	94	36.384	8 <i>.777</i>	88.873	1.00	28.74
		ō	PRO B	94	36.562	8.176	87.808	1.00	25.77
MOTA	3774					9.553	89.112	1.00	
MOTA	3775	N	VAL B	95	35.332				
MOTA	3776	CA	VAL B	95	34.317	9.812	88.103	1.00	
ATOM	3777	CB	VAL B	95	33.035	10.393	88.742	1.00	
	3778	CG1		95	32.067	10.855	87.662	1.00	26.34
MOTA					_	9.346	89.622	1.00	
atom	3779	CG2	VAL B	95	32.378		87.175	1.00	
ATOM	3780	С	VAL B	95	34.912	10.861			
ATOM	3781	0	VAL B	95	35.564	11.793	87.641	1.00	
	3782	N	SER B	96	34.708	10.699	85.871	1.00	28.02
ATOM			SER B	96	35.199	11.647	84.868	1.00	
MOTA	3783	CA				11.705	.84.850	1.00	
ATOM	3784	CB	SER B	96	36.729		84.229	1.00	23.00
ATOM	3785	OG	SER B	96	37.274	10.548	04.229	T.00	23.99

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ATOM	3786	C	SER B	96	34	726	11.127	83.519	1.00 26.22
ATOM	3787		SER B	96	33	943	10.174	83.462	1.00 23.57
	3788		TYR B	97		195	11.744	82.438	1.00 22.83
MOTA				97		818	11.279	81.110	1.00 28.59
MOTA	3789								
MOTA	3790			97		536	12.452	80.165	1.00 31.45
ATOM	3791	CG		97		279	13.203	80.548	1.00 35.09
ATOM	3792	CD	1 TYR B	97	33.	316	14.239	81.480	1.00 32.87
ATOM	3793	CE	1 TYR B	97	32.	148	14.863	81.911	1.00 37.73
ATOM	3794	CD	2 TYR B	97	32.	036	12.812	80.049	1.00 34.85
MOTA	3795	CE		97		858	13.430	80.475	1.00 38.61
	3796	cz		97		924	14.453	81.408	1.00 39.45
ATOM	-			97		768	15.047	81.852	1.00 35.36
ATOM	3797	ОĤ							
ATOM	3798	С	TYR B	97		883	10.354	80.534	1.00 28.93
ATOM	3799	0	TYR B	97		859	9.992	79.358	1.00 28.26
MOTA	3800	N	ALA B	98		822	9.968	81.385	1.00 29.09
ATOM	3801	CA	ALA B	98	37.	866	9.044	80.980	1.00 26.88
ATOM	3802	CB	ALA B	98	39.	167	9.369	81.692	1.00 27.99
ATOM	3803	C	ALA B	98	37.	395	7.657	81.382	1.00 22.53
ATOM	3804	ŏ	ALA B	98		721	6.675	80.722	1.00 21.98
	3805	N	MET B	99		603	7.595	82.453	1.00 23.51
MOTA	_			99			6.326	82.986	1.00 26.36
MOTA	3806	CA	MET B			106			
ATOM	3807	СВ	MET B	99		179	6.568	84.185	1.00 24.05
MOTA	3808	CG	MET B	99		822	7.188	83.875	1.00 28.37
ATOM	3809	SD	MET B	99		966	7.704	85.406	1.00 27.91
MOTA	3810	CE	MET B	99	33.	106	6.227	86.409	1.00 22.12
ATOM	3811	С	MET B	99	35.	430	5.435	81.953	1.00 25.76
ATOM	3812	0	MET B	99	35.	544	4.212	82.031	1.00 26.11
ATOM	3813	N	PHE B	100		724	6.027	80.992	1.00 22.17
ATOM	3814	CA		100		107	5.222	79.940	1.00 22.35
	3815	СВ		100		582	5.133	80.088	1.00 22.01
ATOM		CG	PHE B			947	4.254	79.038	1.00 24.22
ATOM	3816						2.872	79.061	1.00 26.61
MOTA	3817	CD1		100		143			
MOTA	3818	CD2				280	4.813	77.953	1.00 21.22
ATOM	3819	CE1		100		691	2.059	78.012	1.00 26.91
MOTA	3820	CE2	PHE B	100	30.	825	4.010	76.894	1.00 24.80
ATOM	3821	CZ	PHE B	100	31.	033	2.632	76.924	1.00 24.85
ATOM	3822	С	PHE B	100	34.	425	5.695	78.514	1.00 24.86
MOTA	3823	0	PHE B	100	34.	922	4.920	77.694	1.00 21.40
MOTA	3824	N		101	34.	131	6.957	78.204	1.00 24.24
ATOM	3825	CA		101	34.		7.469	76.854	1.00 24.54
	3826	СВ		101	33.		8.926	76.708	1.00 24.46
MOTA	3827	OG1		101	32.		8.985	76.953	1.00 27.64
MOTA				101	34.		9.445	75.297	1.00 22.19
MOTA	3828	CG2						76.483	1.00 25.26
ATOM	3829	С		101	35.		7.387		
MOTA	3830	0		101	36.		6.856	75.430	1.00 25.47
MOTA	3831	N		102	36.		7.916	77.350	1.00 23.74
ATOM	3832	CA	GLY B	102	38.	153	7.867	77.096	1.00 24.53
ATOM	3833	С	GLY B	102	38.	657	6.434	77.046	1.00 24.06
ATOM	3834	0	GLY B	102	39.	346	6.045	76.100	1.00 22.53
ATOM	3835	N	SER B	103	38.	316	5.651	78.067	1.00 22.02
ATOM	3836	CA	SER B		38.		4.253	78.146	1.00 20.45
					38.		3.613	79.427	1.00 25.21
ATOM	3837	CB	SER B		38.		4.166	80.567	1.00 26.48
ATOM	3838	OG	SER B		_			76.938	1.00 20.53
ATOM	3839	С	SER B		38.		3.446		
MOTA	3840	0	SER B		39.		2.669	76.372	1.00 16.82
ATOM	3841	N	SER B	104	37.	014	3.642	76.542	1.00 17.11
ATOM	3842	CA	SER B	104	36.	462	2.937	75.395	1.00 23.32
ATOM	3843	CB	SER B		34.	980	3.289	75.228	1.00 22.93
ATOM	3844	OG	SER B		34.		2.557	74.161	1.00 24.75
	3845	c	SER B		37.		3.288	74.116	1.00 21.97
MOTA		ō	SER B		37.		2.434	73.256	1.00 22.83
MOTA	3846		LEU B		37.		4.549	73.997	1.00 23.00
MOTA	3847	N					5.007	72.825	1.00 25.12
ATOM	3848	CA	LEU B		38.				1.00 29.25
MOTA	3849	CB	LEU B	102	38.		6.536	72.859	
MOTA	3850	CG	LEU B	105	38.		7.289	71.553	1.00 34.27
MOTA	3851	CD1	LEU B	105	37.	662	6.888	70.512	1.00 31.51

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ATOM	3852	CD:	2. LEU E	105	38.629	8.802	71.819	1.00	34.30
	3853	C.	LEU E		39.755	4.374	72.813	1.00	
ATOM									
MOTA	3854	0	LEU F	3 105	40.262	3.955	71.765	1.00	
ATOM	3855	N	ALA E	3 106	40.371	4.293	73.988	1.00	23.62
ATOM	3856	CA	ALA E	106	41.704	3.692	74.115	1.00	22.73
		CB	ALA E		42.263	3.939	75.529		17.46
MOTA	3857								
ATOM	3858	С	ALA E		41.639	2.189	73.846		22.77
ATOM	3859	0	ALA E	106	42.583	1.597	73.316	1.00	24.98
ATOM '	3860	N	THR E	107	40.523	1.567	74.224	1.00	23.54
			THR E		40.355	0.132	74.033		19.76
MOTA	3861	CA							
MOTA	3862	CB	THR E	107	39.236	-0.410	74.947		22.85
MOTA	3863	OG:	1 THR E	107	39.572	-0.128	76.306	1.00	16.29
ATOM	3864	CG2	2 THR E	107	39.085	-1.917	74.787	1.00	17.21
			THR E		40.036	-0.169	72.571		23.47
MOTA	3865	C							19.25
MOTA	3866	0	THR E		40.540	-1.138	72.001		
MOTA	3867	N	GLY E	108	39.191	0.656	71.959		24.87
ATOM	3868	CA	GLY E	108	38.879	0.434	70.560	1.00	21.32
	3869	C	GLY E		40.161	0.594	69.757	1.00	22.01
MOTA						-0.099	68.761		20.46
ATOM	3870	0	GLY E		40.388				
MOTA	3871	N	SER E	109	41.018	1.508	70.197		19.89
MOTA	3872	CA	SER E	109	42.274	1.749	69.499	1.00	19.37
	3873	CB	SER B		42.993	2.956	70.107	1.00	18.28
MOTA			SER B		42.250	4.145	69.882		21.29
MOTA	3874	OG							
MOTA	3875	С	SER B		43.168	0.513	69.542		20.70
MOTA	3876	0	SER B	109	43.940	0.261	68.617	1.00	20.69
ATOM	3877	N	THR B	110	43.065	-0.259	70.616	1.00	20.54
	3878	CA	THR B		43.858	-1.475	70.729	1.00	19.98
MOTA					43.826	-2.043	72.158	1.00	
MOTA	3879	CB	THR B						
MOTA	3880	OG1	THR B		44.632	-1.215	73.007		20.72
ATOM	3881	CG2	THR B	110	44.371	-3.470	72.188	1.00	20.05
ATOM	3882	С	THR B	110	43.333	-2.507	69.738	1.00	21.61
	3883	ō	THR B		44.115	-3.239	69.127		18.11
MOTA							69.567		18.29
MOTA	3884	N	VAL B		42.012	-2.557			
ATOM	3885	CA	VAL B	. 111	41.432	-3:486	68.608		20.36
ATOM	3886	CB	VAL B	111	39.886	-3.494	68.677	1.00	23.94
	3887	CG1			39.324	-4.442	67.619	1.00	24.37
MOTA					39.426	-3.937	70.063		21.60
MOTA	3888	CG2							
MOTA	3889	С	VAL B		41.872	-3.080	67.197		20.35
ATOM	3890	0	VAL B	111	42.146	-3.936	66.362		23.29
ATOM	3891	N	GLN B	112	41.953	-1.775	66.937	1.00	22.07
	3892	CA	GLN B		42.367	-1.290	65.617	1.00	22.34
MOTA					42.199	0.230	65.513		24.54
MOTA	3893	CB	GLN B						
ATOM	3894	CG	GLN B		40.810	0.729	65.843	1.00	
MOTA	3895	CD	GLN B	112	40.700	2.236	65.742		21.19
ATOM	3896	OE1	GLN B	112	40.664	2.794	64.645	1.00	26.73
	3897	NE2			40.667	2.905	66.886	1.00	18.73
ATOM						-1.635	65.363	1.00	
MOTA	3898	C	GLN B		43.826				
ATOM	3899	0	GLN B	112	44.195	-2.020	64.257		15.79
MOTA	3900	N	ALA B	113	44.660	-1.476	66.389	1.00	20.60
	3901	CA	ALA B	113	46.070	-1.790	66.249	1.00	18.02
ATOM		CB	ALA B		46.794	-1.536	67.548	1.00	20.84
MOTA	3902								
MOTA	3903	С	ALA B		46.170	-3.262	65.863		23.78
MOTA	3904	9	ALA B		46.982	-3.642	65.023		19.83
ATOM	. 3905	N	ILE B	114	45.331	-4.091	66.477	1.00	21.45
		CA	ILE B		45.344	-5.511	66.168	1.00	24.26
ATOM	3906						67.191		20.72
ATOM	3907	CB	ILE B		44.507	-6.306			
ATOM	3908	CG2	ILE B		44.476	-7.779	66.800		21.92
ATOM	3909	CG1	ILE B	114	45.116	-6.144	68.593		24.32
		CD1			44.364	-6.872	69.694	1.00	19.01
ATOM	3910						54.753		26.75
ATOM	3911	С	ILE B		44.808	-5.765			
ATOM	3912	0	ILE B		45.305	-6.640	64.032	1.00	20.18
ATOM	3913	N	GLU B	115	43.792	-5.009	64.347		24.69
	3914	CA	GLU B	115	43.243	-5.198	63.005	1.00	29.26
ATOM			GLU B	115	42.043	-4.278	62.770		29.07
atom	3915	CB	G110 D	115			63.800	1 00	32.31
ATCM	3916 ·		GLU B	772	40.940	-4.421		1.00	20 11
ATOM	3917	CD	GLU B	115	39.757	-3.519	63.516	1.00	38.14

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				22 222				40 60
MOTA	3918	OE.	1 GLU B 115	39.980	-2.374	63.072	1.00	40.63
MOTA	3919	OE:	2 GLU B 115	38.607	-3.940	63.758	1.00	39.86
ATOM	3920	C	GLU B 115	44.334	-4.906	61.974	1.00	31.52
ATOM	3921	0	GLU B 115	44.444	-5.603	60.964	1.00	26.43
ATOM	3922	N	GLU B 116	45.141	-3.879	62.234	1.00	26.04
	3923	CA	GLU B 116	46.226	-3.522	61.324		26.21
MOTA								
MOTA	392,4	CB	GLU B 116	46.909	-2.227	61.775	1.00	23.21
ATOM	3925	CG	GLU B 116	46.055	-0.983	61.601	1.00	23.82
				45.576			1.00	
MOTA	3926	CD	GLU B.116		-0.817	60.163		31.43
ATOM	3927	OE:	L GLU B 116	46.425	-0.734	59.253	1.00	26.45
MOTA	3928	OE2		- 44.349	-0.771	59.945	1.00	24.59
MOTA	3929	С	GLU B 116	47.256	-4.644	61.243		28.60
MOTA	3930	0	GLU B 116	47.857	-4.884	60.189	1.00	25.01
MOTA	3931	N	PHE B 117	47.470	-5.324	62.363	1.00	26,22
				48.421				28.05
MOTA	3932	CA	PHE B 117		-6.425	62.400		
MOTA	3933	CB	PHE B 117	48.516	-7.007	63.805	1.00	32.15
ATOM	3934	ĊĠ	PHE B 117	49.278	-8.299	63.869	1.00	33.88
MOTA	3935	CD1		50.656	-8.321	63.713		33.52
ATOM	3936	CD2	PHE B 117	48.604	-9.502	64.054	1.00	32.83
ATOM	3937	CE1	PHE B 117	51.356	-9.521	63.740	1.00	31.67
					-10.710	64.082		35.69
MOTA	3938	CE2						
MOTA	3939	CZ	PHE B 117	50.674	~10.717	63.926	1.00	36.72
MOTA	3940	C	PHE B 117	47.929	-7.508	61.456	1.00	26.43
	3941			48.689	-8.061	60.669		27.61
MOTA		0	PHE B 117					
MOTA	3942	N	LEU B 118	46.642	-7.809	61.551		23.59
ATOM	3943	CA	LEU B 118	46.048	-8.820	60.705	1.00	29.15
	3944		LEU B 118	44.585	-9.039	61.099		28.78
MOTA		CB						
ATOM	3945	CG	LEU B 118	44.375	-9.478	62.557	1.00	35.24
ATOM	3946	CD1	LEU B 118	42.898	-9.763	62.788	1.00	31.92
ATOM	3947	CD2			-10.723	62.856	1 00	33.40
MOTA	3948	С	LEU B 118	46.153	-8.422	59.236		30.15
MOTA	3949	0	LEU B 118	46.350	-9.276	58.379	1.00	27.04
ATOM	3950	N	LYS B 119	46.035	-7.128	58.947	1 00	27.96
ATOM	3951	CA	LYS B 119	46.127	-6.663	57.569		26.69
ATOM	3952	CB	LYS B 119	45.470	-5.291	57.412	1.00	23.94
MOTA	3953	CG	LYS B 119	43.998	-5.260	57.795	1.00	24.41
						57.350		27.53
ATOM	3954	CD	LYS B 119	43.327	-3.970			
MOTA	3955	CE	LYS B 119	44.024	-2.739	57.886	1.00	33.13
ATOM	3956	NZ	LYS B 119	43.371	-1.479	57.428	1.00	27.75
_		C	LYS B 119	47.577	-6.598	57.101		29.12
MOTA	3957							
MOTA	3958	0	LYS B 119	47.864	-6.160	55.984		35.25
ATOM	3959	N	GLY B 120	48.493	-7.034	57.958	1.00	30.25
MOTA	3960	CA	GLY B 120	49.896	-7.037	57.585	1.00	28.38
ATOM	3961	C	GLY B 120	50.642	-5.751	57.861		27.91
MOTA	3962	0	GLY B 120	51.775	-5.582·	57.403	1.00	22.25
ATOM	3963	N	ASN B :21	50.024	-4.836	58.600	1.00	25.42
					-3.587	58.919		29.49
MOTA	3964	CA	ASN B .21	50.695	•			
MOTA	3965	CB	ASN B 121	49.758	-2.389	58.727	1.00	30.07
ATOM	3966	CG	ASN B 121	49.201	-2.307	57.325	1.00	32.25
				49.924	-2.491	56.350		35.44
MOTA	3967							
MOTA	3968	ND2	ASN B 121	47.917	-2.006	57.217	1.00	32.26
ATOM	3969	С	ASN B 121	51.172	-3.637	60.361	1.00	30.92
				50.971	-4.631	61.059		27.08
MOTA	3970	0	ASN B 121					
ATOM	3971	N	VAL B 122	51.810	-2.560 [.]	60.796		28.46
ATOM	3972	CA	VAL B 122	52.309	-2.457	62.155	1.00	29.48
	3973	CB	VAL B 122	53.840	-2.352	62.177		33.40
MOTA								
MOTA	3974	CG1	VAL B 122	54.334	-2.294	63.611		32.22
ATOM	3975	CG2	VAL B 122	54.446	-3.544	61.458	1.00	33.57
	3976	c	VAL B 122	51.713	-1.196	62.748	1.00	29.04
ATOM						62.153		27.47
ATOM	3977	0	VAL B 122	51.800	-0.118			
MOTA	3978	N	ALA B 123	51.100	-1.326	63.918		26.71
	3979	CA	ALA B 123	50.477	-0.177	64.559	1.00	25.62
MOTA						64.447		21.65
MOTA	3980	CB	ALA B 123	48.963	-0.281			
ATOM	3981	С	ALA B 123	50.872	-0.005	66.017	1.00	28.62
	3982	ō	ALA B 123	51.227	-0.965	66.712	1.00	26.96
ATOM				50.805	1.239	66.472	1.00	22.85
MOTA	3983	N	PHE B 124	30.603	1.633	55.472		-4.07

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	2004	CA	PHE B 12	24	51.122	1.577	67.847	1.00 17.31
ATOM	3984	CB	PHE B 12		52.419	2.404	67.876	1.00 16.88
MOTA	3985		PHE B 12		52.762	3.000	69.225	1.00 18.52
ATOM	3986	CG			52.533	2.304	70.403	1.00 17.52
MOTA	3987		PHE B 13			4.245	69.297	1.00 17.88
ATOM	3988	CD2	PHE B 1		53.382		71.638	1.00 25.77
ATOM-	3989	CE1	PHE B 12		52.914	2.837		1.00 21.97
ATOM	3990	CE2	PHE B 12		53.769	4.790	70.517	1.00 21.57
MOTA	3991	CZ	PHE B 1	24	53.535	4.084	71.698	1.00 20.16
	3992	С	PHE B 1	24	49.937	2.348	68.421	1.00 18.77
MOTA	3993	ŏ.	PHE B 1	24	49.462	3.311	67.820	1.00 16.62
ATOM	3994	N	ASN B 1	25	49.418	1.868	69.546	1.00 16.69
MOTA			ASN B 1	25	48.320	2.528	70.238	1.00 16.22
MOTA	3995	CA	ASN B 1	25 .	47.129	1.603	70:435	1.00 12.71
MOTA	3996	CB	ASN B 1		46.095	2.209	71.346	1.00 19.79
MOTA	3997	CG	ASN D I	2J 25	45.930	3.430	71.372	1.00 20.83
MOTA	3998			25 25	45.376	1.371	72.087	1.00 12.31
MOTA	3999	ND2	ASN B 1	2 5		3.004	71.600	1.00 19.19
ATOM	4000	С	ASN B 1	25	48.790		72.585	1.00 20.99
MOTA	4001	0	ASN B 1		48.687	2.280		1.00 19.02
ATOM	4002	N	PRO B 1		49.335	4.226	71.668	1.00 21.39
ATOM	4003	CD	PRO B 1	26	49.595	5.156	70.555	
ATOM	4004	CA	PRO B 1	26	49.833	4.805	72.917	1.00 21.60
	4005	CB	PRO B 1		50.398	6.161	72.459	1.00 21.07
MOTA	4006	CG	PRO B 1	26	49.530	6.487	71.269	1.00 17.70
MOTA		c	PRO B 1	26	48.808	4.942	74.034	1.00 20.69
ATOM	4007		PRO B 1		49.178	5.053	75.198	1.00 19.79
MOTA	4008	0			47.525	4.937	73.689	1.00 16.67
ATOM	4009	N	ALA B 1	27	46.476	5.065	74.698	1.00 20.44
ATOM	4010	CA	ALA B 1			5.609	74.066	1.00 19.56
ATOM	4011	CB	ALA B 1		45.198	3.747	75.401	1.00 20.80
ATOM	4012	С	ALA B 1		46.169	3.742	76.472	1.00 19.47
ATOM	4013	0	ALA B 1	.27	45.555			1.00 20.52
ATOM	4014	N	GLY B 1	.28	46.587	2.634	74.800	1.00 19.43
ATOM	4015	CA	GLY B 1	.28	46.325	1.333	75.399	1.00 20.56
ATOM	4016	С	GLY B 1	.28	47.327	0.910	76.463	1.00 20.30
ATOM	4017	0	GLY B 1	.28 .	48.182	1.697	76.869	1.00 18.37
ATOM	4018	N	GLY B 1	.29	47.215	-0.333	76.929	1.00 19.68
MOTA	4019	CA	GLY B 1	.29	48.136	-0.820	77.943	1.00 19.93
	4020	C	GLY B 1	.29	47.620	-0.619	79.358	1.00 25.25
ATOM	4021	ō	GLY B 1		48.383	-0.686	80.329	1.00 18.98
ATOM	4022	N	MET B 1	30	46.317	-0.374	79.474	1.00 16.04
ATOM		CA	MET B 1		45.677	-0.161	80.768	1.00 19.26
ATOM	4023		MET B 1		44.301	0.451	80.519	1.00 17.94
ATOM	4024	CB	MET B 1		44.413	1.728	79.653	1.00 22.95
ATOM	4025	CG			42.873	2.615	79.307	1.00 31.83
ATOM	4026	SD	MET B 1		41.957	1.358	78.382	1.00 20.22
ATOM	4027	CE	MET B 1			-1.548	81.421	1.00 22.63
ATOM	4028	C	MET B 1	130	45.598		81.486	1.00 16.24
ATOM	4029	0	MET B 1	L30	44.546	-2.173	81.932	1.00 18.42
ATOM	4030	N	HIS B 1		46.737	-1.999	82.472	1.00 17.07
ATOM	4031	CA	HIS B 1		46.853	-3.343		1.00 17.61
TOM	4032	CB	HIS B 1	131	48.323	-3.804	82.341	1.00 14.01
TOM	4033	CG	HIS B 1	L31	49.316	-2.979	83.106	
	4034	CD2	HIS B 1	131	49.138	-1.904	83.915	1.00 13.47
ATOM	4035	ND1	HIS B 1	131	50.680	-3.190	83.051	1.00 18.00
ATOM		CF1	HIS B 1	131	51.297	-2.281	83.789	1.00 15.27
ATOM	4036	NE2		131	50.384	-1.489	84.324	1.00 17.21
ATOM	4037		HIS B 1	131	46.329	-3.724	83.852	1.00 16.41
MOTA	4038	C	TIS B I	121	46.452	-4.883	84.236	1.00 19.37
atom	4039	0	HIS B 1	131	45.721	-2.794	84.586	1.00 18.64
ATOM:	4040	N	HIS B 1	132		-3.112	85.936	1.00 20.87
ATOM	4041	CA	HIS B 1	132	45.241	_1 035	86.885	
MOTA	4042	CB	HIS B 1	132	45.513	-1.935	87.152	1.00 20.00
ATOM	4043	CG	HIS B 1	132	46.966	-1.686		
ATOM	4044	CD2	HIS B 1	132	47.715	-0.563	87.030	
ATOM	4045	ND1	HIS B I	132	47.810	-2.655	87.659	
	4046	CEI	HIS B	132	49.014	-2.139	87.837	
ATOM		MES	HIS B	132	48.984	-0.872	87.462	1.00 14.88
ATOM	404.7		HIS B	132	43.778	-3.547	86.136	1.00 22.83
-TOM	4048	С	HIS B	132	43.478	-4.298	87.076	1.00 17.84
LTOM	4049	0	HIP B		23.2.0			

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							05 051	1 00 15 51
MOTA	4050	N	ALA E	3 133	42.878	-3.088	85.271	1.00 16.54
	4051	CA	ALA E	3 133	41.457	-3.396	85.424	1.00 19.13
MOTA								
MOTA	4052	CB	ALA E	3 133	40.654	-2.704	84.328	1.00 23.56
ATOM	4053	С	ALA E	3 133	41.127	-4.883	85.439	1.00 23.12
						-5.677	84.696	1.00 18.03
ATOM	4054	0	ALA E		41.718			
ATOM	4055	N	PHE E	3 134	40.181	-5.257	86.294	1.00 19.69
			PHE E		39.762	-6.649	86.365	1.00 19.35
ATOM	4056	CA						
MOTA	4057	CB	PHE E	134	39.583	-7.122	87.818	1.00 21.26
	4058	CG	PHE E	134	40.837	-7.053	88.646	1.00 23.41
ATOM								
ATOM	4059	CDT	PHE E	134	41.041	-6.009	89.544	1.00 24.25
MOTA	4060	CD2	PHE E	134	41.820	-8.027	88.522	1.00 22.80
					42.207	-5.935	90.311	1.00 23.36
MOTA	4061	CEL						
ATOM	4062	CE2	PHE B	134	42.997	-7.964	89.283	1.00 27.74
	4063	CZ	PHE E	134	43.190	-6.917	90.178	1.00 24.05
MOTA								
ATOM	4064	С	PHE B	134	38.444	-6.816	85.621	1.00 18.60
ATOM	4065	0	PHE B	134	37.815	-5.849	85.196	1.00 13.82
					38.050	-8.064	85.454	1.00 19.78
MOTA	4066	N	LYS E					
ATOM	4067	CA	LYS B	135	36.813	-8.421	84.782	1.00 28.09
	4068	CB	LYS B		36.501	-9.879	85.125	1.00 34.06
MOTA								•
MOTA	4069	CG	LYS B			-10.310	84.953	1.00 42.76
MOTA	4070	CD	LYS B	135	34.927	-11.745	85.437	1.00 48.44
	4071		LYS B			-12.152	85.531	1.00 55.66
MOTA		CE						
ATOM	4072	NZ	LYS B			-11.332	86.544	1.00 51.65
ATOM	4073	С	LYS B	135	35.639	-7.512	85.172	1.00 28.27
			LYS B		34.927	-6.999	84.309	1.00 24.86
MOTA	4074	0						
MOTA	4075	N	SER B	136	35.450	-7.292	86.470	1.00 29.89
MOTA	4076	CA	SER B	136	34.331	-6.477	86.933	1.00 30.86
	4077	CB	SER B		33.282	-7.388	87.582	1.00 31.57
MOTA								
MOTA	4078	OG	SER B	136	32.916	-8.434	86.698	1.00 45.10
MOTA	4079	С	SER B	136	34.705	-5.380	87.923	1.00 31.50
	4080	0	SER B		33.887	-4.997	88.765	1.00 24.54
ATOM								
MOTA	4081	N	ARG B		35.920	-4.854	87.835	1.00 22.63
ATOM	4082	CA	ARG B	137	36.291	-3.826	88.794	1.00 25.51
	4083	CB	ARG B	137	36.629	-4:486	90.136	1.00 29.62
MOTA							91.318	1.00 36.21
MOTA	4084	CG	ARG B		36.391	-3.578		
ATOM	4085	CD	ARG B	137	36.874	-4.160	92.631	1.00 40.79
	4086	NE	ARG B	137	36.365	-3.357	93.744	1.00 45.95
ATOM						-3.369	94.973	1.00 41.97
MOTA	4087	cz	ARG B		36.863			
ATOM	4088	NH1	ARG B	137	37.897	-4.144	95.263	1.00 43.42
	4089	NH2	ARG B	137	36.322	-2.604	95.913	1.00 46.65
MOTA					37.461	-2.956	88.339	1.00 24.73
MOTA	4090	С	ARG B					
MOTA	4091	0	ARG B	137	38.420	-3.441	87.734	1.00 19.32
	4092	N	ALA B	138	37.372	-1.663	88.631	1.00 16.77
MOTA						-0.733	88.270	1.00 18.50
MOTA	4093	CA	ALA B		38.428			
MOTA	4094	CB	ALA B	138	37.939	0.694	88.401	1.00 17.24
	4095	C	ALA B	138	39.597	-0.964	89.216	1.00 22.62
MOTA							90.346	1.00 18.98
MOTA	4096	0	ALA B		39 411	-1.419		
MOTA	4097	N	ASN B	139	40.301	-0.641	88.759	1.00 20.82
	4098	CA	ASN B	139	41.989	-0.828	89.585	1.00 25.17
ATOM							89.689	1.00 20.59
ATOM	4099	CB	ASN B		42.311	-2.329		
MOTA	4100	CG	ASN B	139	43.556	-2.608	90.511	1.00 27.70
	4101		ASN B	139	43.726	-2.057	91.592	1.00 22.43
MOTA							90.010	1.00 24.43
MOTA	4102	NDZ	ASN B		44.420	-3.487		1.00 24.45
MOTA	4103	С	ASN B	139	43.176	-0.062	89.020	1.00 22.37
	4104	0	ASN B		43.338	0.038	87.799	1.00 17.50
MOTA							89.920	1.00 21.67
MOTA	4105	И	GLY B		43.984	0.496		
MOTA	4106	CA	GLY B	140	45.166	1.249	89.524	1.00 23.06
	4107		GLY B		45.005	2.268	88.402	1.00 26.29
MOTA						2.301	87.479	1.00 22.47
ATOM	4108	0	GLY B		45.827			1.00 44.37
MOTA	4109	N	PHE B	141	43.958	3.093	88.473	1.00 22.33
	4110	CA	PHE B	141	43.694	4.126	87.461	1.00 19.01
ATOM		C71	DITE E	1/1	44.996	4.806	86.997	1.00 22.90
MOTA	4111	CB	PHE B	141				1.00 22.70
ATOM	4112	CG	PHE B	141	45.810	5.433	88.097	1.00 23.17
	4113	CD1	PHE B	141	47.114	5.851	87.842	1.00 22.17
ATOM ·		CDO	PARE E	1 4 1	45.281	5.635	.89.366	1.00 23.40
ATOM	4114	حريب	PHE B	441				1.00 24 02
ATOM	4115	CEl	PHE B	141	47.876	6.462	88.833	1.00 24.02
WION								

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						c 244	90.361	1.00 23.03
ATOM	4116	CE2	PHE B	141	46.033	6.244		1.00 25.15
MOTA	4117	CZ	PHE B	141	47.335	6.658	90.092	
ATOM	4118	С	PHE B	141	43.029	3.538	86.214	1.00 23.69
ATOM	4119	Ō	PHE B	141	42.596	4.283	85.335	1.00 18.88
	4120	N	CYS B		42.962	2.211	86.122	1.00 15.03
MOTA					42.380	1.578	84.938	1.00 19.55
MOTA	4121	CA	CYS B				84.552	1.00 20.38
ATOM	4122	CB	CYS B		43.193	0.336		1.00 20.30
MOTA	4123	SG	CYS B	142	44.933	0.662	84.190	
MOTA	4124	С	CYS B	142	40.923	1.171	85.098	1.00 22.77
ATOM	4125	Ö	CYS B	142	40.561	0.514	86.082	1.00 23.04
	4126	N	TYR B		40.094	1.557	84.130	1.00 15.24
MOTA			TYR B		38.675	1.194	84.155	1.00 21.97
ATOM	4127	CA				2.372	83.723	1.00 18.06
MOTA	4128	CB	TYR B		37.795		84.535	1.00 24.34
MOTA	4129	CG	TYR B		38.016	3.622	04.014	
ATOM	4130	CD1	TYR B		39.038	4.516	84.214	1.00 23.20
ATOM	4131	CE1	TYR B	143	39.265	5.658	84.991	1.00 27.42
	4132	CD2	TYR B		37.226	3.892	85.652	1.00 19.15
MOTA		CE2	TYR B		37.441	5.023	86.432	1.00 21.92
ATOM	4133				38.458	5.900	86.099	1.00 23.94
MOTA	4134	CZ	TYR B			7.015	86.877	1.00 22.37
MOTA	4135	OH	TYR B	143	38.655		83.218	1.00 19.91
ATOM	4136	C	TYR B		38.431	0.008		
MOTA	4137	0	TYR B	143	37.665	-0.902	83.535	1.00 22.50
ATOM	4138	N	ILE B	144	39.083	0.026	82.061	1.00 19.20
	4139	CA	ILE B	144	38.938	-1.055	81.082	1.00 19.68
ATOM	4140	CB		144	38.282	-0.528	79.787	1.00 20.26
ATOM			ILE B		38:151	-1.649	78.760	1.00 15.37
ATOM	4141	CG2			36.901	0.053	80.113	1.00 20.93
ATOM	4142	CG1	ILE B				78.917	1.00 23.75
MOTA	4143	CD1	ILE B		36.198	0.697		1.00 22.78
MOTA	4144	С	ILE B		40.320	-1.627	80.774	
MOTA	4145	0	ILE B	144	41.281	-0.873	80.600	1.00 22.01
ATOM	4146	N	ASN B	145	40.422	-2.956	80.723	1.00 23.18
	4147	CA	ASN B	145	41.698	-3.623	80.451	1.00 20.63
MOTA	4148	CB	ASN B		41.778	-4.935	81.243	1.00 17.81
MOTA	-			145	43.188	-5.531	81.268	1.00 25.17
MOTA	4149	CG				-5.742	80.227	1.00 23.63
MOTA	4150		ASN B	145	43.804			1.00 22.69
MOTA	4151	ND2	ASN B	145	43.693	-5.819	82.472	1.00 21.18
MOTA	4152	С	ASN B	145	41.780	-3.918	78.955	
ATOM	4153	0	ASN B	145	41.389	-5.002	78.508	1.00 17.80
	4154	N	ASN B	146	42.293	-2.968	78.177	1.00 15.23
MOTA	4155	CA	ASN B	146	42.367	-3.175	76.733	1.00 19.71
ATOM		CB	ASN B	146	42.773	-1.880	76.015	1.00 17.65
ATOM	4156			146	44.196	-1.458	76.306	1.00 19.86
ATOM	4157	CG	ASN B			-1.735	75.532	1.00 20.27
ATOM	4158		ASN B	146	45.109		77.435	1.00 11.85
ATOM	4159	ND2	ASN B	146	44.395	-0.798		1.00 19.07
MOTA	4160	С	asn b	146	43.277	-4.342	76.331	
MOTA	4161	0	ASN B	146	43.030	-4.996	75.328	1.00 18.61
ATOM	4162	N	PRO B	147	44.358	-4.598	77.082	1.00 17.78
	4163	CD	PRO B	147	44.953	-3.919	78.240	1.00 18.13
MOTA			PRO B	147	45.197	-5.735	76.678	1.00 19.98
ATOM	4164	CA			46.338	-5.694	77.698	1.00 24.29
MOTA	4165	CB	PRO B				78.020	1.00 26.27
ATOM	4166	CG	PRO B	14/	46.425	-4.201	76.020	1.00 20.91
MOTA	4167	С	PRO B	147	44.377	-7.041	76.757	1.00 20.51
MOTA	4168	0	PRO B	147	44.461	-7.892	75.871	1.00 17.58
	4169	N	ALA B	148	43.568	-7.172	.77.809	1.00 15.81
MOTA		CA	ALA B	148	42.732	-8.362	78.008	1.00 19.82
MOTA	4170		ALA B		42.049	-8.312	79.372	1.00 17.50
MOTA	4171	CB				-8.473	76.903	1.00 22.58
ATOM	4172	С	ALA B	140	41.683		76.404	1.00 18.38
ATOM	4173	0	ALA B		41.419	-9.567		1.00 22.48
MOTA	4174	N	VAL B		41.080	-7.341	76.540	
ATOM	4175	CA	VAL B		40.086	-7.300	75.466	1.00 19.04
	4176	CB	VAL B		39.503	-5.877	75.281	1.00 18.96
ATOM		CG1		149	38.691	-5.800	73.988	1.00 17.32
MOTA	4177		VAL B		38.621	-5.531	76.462	1.00 15.33
ATOM	4178	CG2	AWD D	1/0	40.763	-7.709	74.166	1.00 22.12
ATOM	4179	С	VAL B	147	40.703	_	73.421	1.00 21.83
ATOM	4180	0	VAL B	149	40.240	-8.535		1.00 19.51
ATOM	4181	N	GLY B	150	41.927	-7.120	73.903	T.00 T3.3T

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		~-	ar 1/ n	1 5 0	42 657	7 422	72 600	3 00	19.32
MOTA	4182	CA	GLY B	150	42.657	-7.433	72.689		
MOTA	4183	С	GLY B	150	43.033	-8.901	72.606	1.00	19.59
					42.862	-9.550	71.568	1 00	22.28
ATOM	4184	0		150					
MOTA	4185	N	ILE B	151	43.558	-9.435	73.700	1.00	19.51
	4186	CA		151	43 958	-10.834	73.723	1 00	23.21
MOTA									
ATOM	4187	CB	ILE B	151	44.666	-11.175	75.053	1.00	23.50
ATOM	4188	CG2	ILE B	151	44.918	-12.679	75.158	1.00	20.01
		_					75.129		21.98
MOTA	4189	CG1	ILE B	121		-10.394			
MOTA	4190	CD1	ILE B	151	46.716	-10.502	76.457	1.00	21.24
						-11.741	73.490	1 00	28.40
MOTA	4191	С	ILE B						
MOTA	4192	0	ILE B	151 .	42.832	-12.692	72.706	1.00	22.96
				152	41 623	-11.450	74.144	7 00	27.32
MOTA	4193	N							
MOTA	4194	CA	GLU B	152	40.417	-12.265	73.939	1.00	27.62
ATOM	4195	CB	GLU B	152	39.294	-11.845	74.886	1.00	26.46
					39.533	-12.200	76.347		28.26
MOTA	4196	CG		152					
MOTA	4197	CD	GLU B	152	39.613	-13.708	76.5 9 2	1.00	31.10
				152	39.668	-14.123	77.767	7 00	29.55
MOTA	4198	OE1							
ATOM	4199	OE2	GLU B	152	39.626	-14.481	75.617	1.00	30.51
ATOM	4200	С	GLU B	152	39.948	-12.125	72.497	1.00	30.30
						-13.082	71.893		25.58
MOTA	4201	0	GLU B						
MOTA	4202	N	TYR B	153	40.093	-10.923	71.948	1.00	26.23
	4203	CA		153	39 720	-10.669	70.563	1.00	28.19
MOTA									
ATOM	4204	CB	TYR B	153	40.082	-9.235	70.190		27.94
MOTA	4205	CG	TYR B	153	39.879	-8.886	68.735	1.00	28.46
				153	38.618	-8.560	68.240	1 00	25.69
MOTA	4206	CD1							
MOTA	4207	CEl	TYR B	153	38.447	-8.195	66.898	1.00	30.73
ATOM	4208	CD2	TYR 3	153	40.962	-8.847	67,856	1.00	24.82
			TYR B		40.801	-8.488	66.526	1 00	29.26
MOTA	4209	CE2							
ATOM	4210	CZ	TYR B	153	39.547	-8.161	66.054		31.25
ATOM	4211	OH	TYR B	153	39.406	-7.803	64.735	1.00	34.22
				153	40.513	-11.627	69.674	1.00	28.11
ATOM	4212	С							
MOTA	4213	0	TYR B	153	39. <i>9</i> 75	-12.248	68.759		22.06
ATOM	4214	N	LEU B	154	41.810	-11.725	69.944	1.00	26.77
	4215	CA		154	42.681	-12.:597	69.168	1.00	28.79
ATOM		•							28.06
MOTA	4216	CB	LEU B		44.142	-12.386	69.592		
MOTA	4217	CG	LEU B	154	44.789	-11.087	69.083	1.00	27.71
ATOM	4218	CD1	LEU B	154	46.119	-10.860	69.759	1.00	34.15
						-11.171	67.571		26.71
ATOM	4219	CD2		154					
ATOM	4220	С	LEU B	154		-14.074	69.274		26.98
ATOM	4221	0	LEU B	154	42.282	-14.787	68.271	1.00	29.88
-				155		-14.536	70.480	1 00	23.19
MOTA	≟222	N							
MOTA	4223	CA	ARG B	155	41.622	-15.936	70.669		29.47
MOTA	4224	CB	ARG B	155	41.339	-16.230	72.144	1.00	28.53
			ARG B		42.527	-15.965	73.053	1.00	35.03
MOTA	4225	CG							
ATOM	4226	CD	ARG B	155	42.212	-16.276	74.507		39.42
ATOM	4227	NE	ARG B	155	42.165	-17.706	74.792	1.00	30.99
						-18.209	75.986	1 00	41.33
MOTA	4228	CZ	ARG B						
MOTA	4229	NHl	ARG B	155	41.591	-17.394	77.002	1.00	38.47
	4230	NH2	ARG B	155	41.872	-19.523	76.178	1.00	40.67
MOTA									29.07
ATOM	4231	C	ARG B	155 .	40.393	-16.260	69.832		
MOTA	4232	0	ARG B	155	40.325	-17.311	69.203	1.00	25.31
	4233	N	LYS B			-15.357	69.828	1.00	28.99
ATOM			113 5	150					34.63
MOTA	4234	CA	LYS B	156		-15.573	69.038		
ATOM	4235	CB	LYS B	156	37.148	~14.534°	69.386	1.00	36.63
	4236	CG	LYS B			-14.883	70.646	1.00	42.18
ATOM									51.38
ATOM	4237	CD	LYS B		37.292	-14.900	71.868		
ATOM	1238	CΞ	LYS B	156	36.685	-15.712	73.009		52.76
		NZ		156		-17.172	72.677	1.00	51.29
ATOM	4239							: 00	34.66
ATOM.	4240	С	LYS B			~15.562	67.538		
ATOM	4241	0	LYS B	156	37.722	-16.088	66.754	1.00	33.53
	4242	N	LYS B			-14.966	67.140	1.00	30.06
atem							65.734		31.36
ATOM	-243	CA		157		-14.945		1.00	22.20
ATCM	1244	CB	LYS B	157	40.888	-13.746	65.418		29.79
	1245	CG	LYS B			-12.426	65.359		31.52
ATOM				_			64.239		28.48
ATOM	4246	CD	LYS B			-12.424			21 00
ATOM	4247	CE	LYS B	157	38.395	-11.101	64.171	1.00	31.90
							-		

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ATOM	4248	NZ	LYS B	157		37.406	-11.080	63.054	1.00 32.91
			LYS B				-16.234	65.381	1.00 31.92
ATOM	4249	C						64.246	1.00 33.58
ATOM	4250	0	LYS B				-16.421		
MOTA	4251	N	GLY B	158		40.890	-17.111	66.368	1.00 28.97
ATOM	4252	CA	GLY B			41.546	-18.379	66.112	1.00 28.98
			GLY B				-18.569	66.622	1.00 33.33
MOTA	4253	С						66.522	1.00 30.58
ATOM	4254	0	GLY B				-19.672		
MOTA	4255	N	PHE B	159		43.578	-17.521	67.164	1.00 32.80
	4256	CA	PHE B	. 159		44.937	-17.657	67.678	1.00 28.89
MOTA			PHE B				-16.286	67.934	1.00 30.33
ATOM	4257	CB -							1.00 28:53
ATOM	4258	CG	PHE B	159			-15.470	66.692	
ATOM	4259	CD1	PHE B	159		44.682	-14.787	66.121	1.00 24.58
	4260	CD2	PHE B	159		46.989	-15.420	66.068	1.00 24.21
ATOM			PHE B			44 849	-14.066	64.948	1.00 25.26
ATOM	4261	CE1					-14.706	64.895	1.00 23.66
MOTA	4262	CE2	PHE B						
ATOM	4263	CZ	PHE B	159			-14.026	64.332	1.00 26.65
ATOM	4264	С	PHE B	159		44.969	-18.484	68.958	1.00 30.92
	4265	ō	PHE B			44.102	-18.334	69.820	1.00 24.26
ATOM			THE D	160		-	-19.347	69.077	1.00 28.86
MOTA	4266	N	LYS B						1.00 30.27
ATOM	4267	CA	LYS B				-20.224	70.237	
MOTA	4268	CB	LYS B	160		46.085	-21.692	69.800	1.00 32.05
	4269	CG	LYS B			44.806	-22.117	69.113	1.00 41.13
MOTA			LYS B				-23.621	68.826	1.00 40.73
ATOM	4270	CD			•			67.904	1.00 43.16
ATOM	4271	CE	LYS B				-24.031		
ATOM	4272	NZ	LYS B	160			-23.408	66.554	1.00 48.69
ATOM	4273	C	LYS B	160		47.394	-19.997	71.048	1.00 28.23
	4274		LYS B			47.552	-20.561	72.130	1.00 25.29
MOTA							-19.206	70.520	1.00 28.51
ATOM	4275	N	ARG B						1.00 25.84
MOTA	4276	CA	ARG B	161			-18.921	71.247	
MOTA	4277	CB	ARG B	161		50.724	-19.719	70.667	1.00 25.33
	4278	CG	ARG B	161	•	50.551	-21.245	70.781	1.00 27.47
ATOM			ARG B				-21.985	70.394	1.00 32.27
MOTA	4279	CD						69.002	1.00 34.90
ATOM	4280	NE	ARG B			52.218	-21.761		
MOTA	4281	CZ	ARG B	161		51.584	-22:276	67.954	1.00 38.45
MOTA	4282	NH1	ARG B	161		50.527	-23.056	68.130	1.00 38.77
	4283	NH2	ARG B			51 999	-22.000	66.725	1.00 38.64
MOTA							-17.421	71.182	1.00 30.40
MOTA	4284	С	ARG B					70.218	1.00 27.50
ATOM	4285	0	ARG B				-16.912		
ATOM	4286	N	ILE B	162		49.376	-16.722	72.221	1.00 25.64
MOTA	4287	CA	ILE B	162		49.515	-15.273	72.303	1.00 27.44
	4288	СВ	ILE B			48.134	-14.618	72.545	1.00 24.53
MOTA							-13.101	72.473	1.00 25.49
ATCM	4289	CG2	ILE B					71.487	1.00 29.46
ATOM	4290	CG1	ILE B				-15.101		
ATOM	4291	CD1	ILE B	162		45.688	-14.707	71.758	1.00 31.94
ATOM	4292	C-	ILE B	162		50.465	-14.868	73.429	1.00 22.68
		ō	ILE B				-15.302	74.568	1.00 24.25
ATOM	4293						-14.042	73.100	1.00 19.49
ATOM	4294	N	LEU B						1.00 17.57
ATOM	4295	CA	LEU B				-13.561	74.081	
ATOM	4296	CB	LEU B	163		53.850	-13.686	73.528	1.00 20.54
	4297	CG	LEU B			54.979	-12.975	74.295	1.00 18.84
ATOM			LEU B				-13.538	75.690	1.00 20.55
MOTA	4298	CD1						73.556	1.00 18.57
ATOM	4299	CD2	LEU B	163		56.293	-13.148		1 00 16 30
ATCM	4300	С	LEU B	163		52.158	-12.099	74.430	1.00 16.20
	4301	o	LEU B			51.898	-11.277	73.549	1.00'16.09
MOTA			TYR B			52 227	-11.780	75.715	1.00 14.05
ATOM	4302	N				52.027	-10.411	76.191	1.00 16.21
ATOM	4303	CA	TYR B						1.00 16.01
ATOM	4304	СЗ	TYR B	164			-10.323	77.070	
ATOM	4305	CG	TYR B	164		50.534	-8.948	77.667	1.00 15.08
	4306	CD1	TYR B			50.148	-7.869	76.869	1.00 19.51
ATOM			TYR 5			49.948	-6.597	77.418	1.00 12.88
ATOM	4307	CE1						79.021	1.00 14.07
ATOM	4308	CD2	TYR B			50.715	-8.724		
ATOM	4309	CE2	TYR B	164		50.520	-7.463	79.583	1.00 13.66
	4310	CZ	TYR B	164		50.139	-6.407	78.782	1.00 14.72
ATOM			TYR B			49.952	-5.163	79.354	1.00 13.54
ATOM	4311	OH					-10.017	77.018	1.00 19.14
ATOM	4312	С	TYR B			33.240	10.017		1.00 26.51
=TOM	4313	0	TYR B	164		55.559	-10.642	78.036	1.00 20.31

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MOTA	4314	N	ILE B	165		53.964	-8.992	76.573	1.00 22.40
MOTA	4315	CA		165		55.148	-8.518	77.285	1.00 17.72 1.00 22.51
MOTA	4316	CB		165		56.352	-8.465 -7.902	76.343 77.079	1.00 16.36
MOTA	4317	CG2	ILE B	165		57.582 56.632	-9.880	75.818	1.00 19.82
MOTA	4318	CG1	ILE B	165 165		57.721	-9.942	74.742	1.00 21.74
MOTA	4319	CD1 C		165		54.851	-7.126	77.850	1.00 22.54
ATOM	4320 4321	0	ILE B	165		54.478	-6.223	77.111	1.00 16.60
MOTA MOTA	4322	N	ASP B	166		55.046	-6.961	79.156	1.00 15.78
MOTA	4323	CA	ASP B	166		54.740	-5.704	79.840	1.00 20.62
ATOM	4324	CB	ASP B	166		53.719	-5.996	80.949 81.486	1.00 17.57 1.00 25.39
MOTA	4325	CG	ASP B	166		53.063 53.779	-4.742 -3.859	82.003	1.00 19.68
ATOM	4326	OD1	ASP B	166 166		51.824	-4.637	81.377	1.00 29.22
MOTA	4327 4328	Ċ	ASP B	166		55.976	-5.002	80.423	1.00 19.01
ATOM ATOM	4329	0	ASP B	166		56.509	-5.412	81.456	1.00 19.74
ATOM	4330	N	LEU B	167		56.414	-3.923	79.775	1.00 17.88
ATOM	4331	CA	LEU B	167		57.598	-3.211	80.235	1.00 14.99 1.00 19.22
MOTA	4332	CB	LEU B	167		58.412	-2.710	79.044 78.069	1.00 22.68
MOTA	4333	CG	LEU B	167		58.871 59.835	-3.799 -3.179	77.074	1.00 25.35
MOTA	4334		LEU B	167 167		59.570	-4.943	78.808	1.00 17.54
MOTA	4335 4336	CD2	LEU B	167		57.284	-2.059	81.183	1.00 17.49
ATOM ATOM	4337	Ö	LEU B	167		58.189	-1.359	81.639	1.00 13.39
ATOM	4338	N	ASP B	168		56.003	-1.878	81.479	1.00 20.03
ATOM	4339	CA	ASP B	168		55.549	-0.848	82.412 82.597	1.00 21.98 1.00 21.21
MOTA	4340	CB	ASP B	168		54.030	-0.955 0.186	83.428	1.00 24.92
ATOM	4341	CG	ASP B	168 168		53.453 56.241	-1.139	83.753	1.00 22.98
MOTA	4342 4343	С 0	ASP B	168		56.447	-2.304	84.091	1.00 18.36
MOTA MOTA	4344		ASP B	168		52.849	1.099	82.825	1.00 22.03
ATOM	4345	OD2	_	168		53.606	0.189	84.676	1.00 18.43
ATOM	4346	N	ALA B	169		56.581	-0.095	84.514	1.00 15.46 1.00 18.73
ATOM	4347	CA	ALA B	169	•	57.263	-0.268 1.084	85.807 86.323	1.00 11.98
ATOM	4348	CB	ALA B	169 169		57.764 56.400	-0.940	86.886	1.00 21.82
ATOM	4349 4350	C O	ALA B	169		56.886	-1.262	87.980	1.00 22.51
MOTA MOTA	4351	N	HIS B	170		55.120	-1.134	86.600	1.00 18.75
ATOM	4352	CA	HIS B	170		54.238	-1.776	87.570	1.00 22.70
MOTA	4353	С	HIS B			53.716	-3.096	87.015 85.809	1.00 22.11 1.00 21.94
MOTA	4354	0	HIS B			53.536 53.050	-3.244 -0.867	87.927	1.00 21.28
MOTA	4355	CB	HIS B	170 170		53.449	0.475	88.460	1.00 18.89
MOTA	4356 4357	CG	HIS B			53.695	1.539	87.626	1.00 19.13
MOTA MOTA	4358	CE1				54.046	2.539	88.412	1.00 19.41
ATOM	4359	CD2	_	170		53.660	0.854	89.746	1.00 19.02 1.00 20.45
ATOM	4360	NE2	HIS B			54.042	2.174	89.710 87.907	1.00 20.45 1.00 19.20
ATOM	4361	N	HIS B			53.474	-4.047 -5.352	87.519	1.00 21.20
ATOM	4362	CA	HIS B		-	52.961 52.964	-6.284	.88.722	1.00 22.00
MOTA	4363	CB	HIS B			52.541	-7.683	88.400	1.00 24.64
MOTA	4364 4365	CD2	HIS B	171		53.056	-8.594	87.540	1.00 19.19
atom Atom	4366	ND1	HIS B	171		51.441	-8.279	88.979	1.00 25.71
MOTA	4367	CE1	HIS B	171		51.295	-9.497	88.487	1.00 25.30 1.00 24.71
ATOM	4368	NE2	HIS B	171		52.261	-9.713	87.612 86.943	1.00 23.91
MOTA	4369	Ç	HIS B	171		51.549	-5.306 -4.620	87.479	1:00 18.93
MOTA	4370	0	HIS B	177		50.677 51.332	-6.062	85.865	1.00 15.36
MOTA	4371	N CA	CYS B	172		50.036	-6.141	85.207	1.00 20.03
MOTA MOTA	4372 4373	CB	CYS B	172		50.240	-6.534	83.732	1.00 22.46
ATOM	4374	SG	CYS B	172		51.259	-8.030	83.419	1.00 23.49
ATOM	4375	c	CYS B	172		49.110	-7.146	85.913 85.327	1.00 18.05 1.00 18.23
ATOM	4376	0	CYS B	172		48.712	-8.151 -6.871	85.327 87.170	1.00 16.78
ATOM	4377	N	ASP B	173		48.767 47.909	-6.871 -7.776	87.928	1.00 18.81
ATOM	4378	CA	ASP B	173.		47.638	-7.236	89.344	1.00 20.39
atem	1379	CB	ADP D	113		-,		•	

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-5.871
                                   46.961
46.564
                                                              1.00 23.40
                                                     89.354
                  ASP B 173
        4380
              CG
ATOM
                                            ~5.435
                                                              1.00 18.64
                                                     90.455
        4381
              OD1
                   ASP
                       B 173
ATOM
                                   46.834
                                            -5.231
                                                     88.291
                                                              1.00 19.24
                       B 173
              OD2
                   ASP
MOTA
        4382
                                                              1.00 17.46
                                   46.595
                                            -8.116
                                                     87.219
MOTA
        4383
              С
                   ASP B 173
                                                              1.00 15.53
                                                     87.224
                   ASP B 173
                                   46.162
                                            -9.272
        4384
              0
ATOM
                                   45.978
                                            -7.130
                                                     86.580
                                                              1.00 13.46
                   GLY B 174
        4385
MOTA
              N
                                   44.733
                                                     85.876
                                                              1.00 18.18
                                            ~7.391
MOTA
        4386
              CA
                   GLY B 174
                                                              1.00 17.85
                                                     84.741
                   GLY B 174
                                   44.904
                                            -8.392
        4387
              С
ATOM
                                            -9.316
                                                     84.583
                                                              1.00 18.27
                   GLY B 174
                                   44.104
ATOM
        4388
              Ω
                                                              1.00 16.14
                                   45.951
                                                     83.943
                   VAL B 175
                                            ~8.214
ATOM
        4389
              N
                                                              1.00 17.00
                                                     82.829
                   VAL B 175
                                   46.206
                                            ~9.111
        4390
              CA
MOTA
                                   47.305
                                            -8.552
                                                     81.902
                                                              1.00 27.22
                   VAL B 175
        4391
              CB
MOTA
                                                     80.731
                                                              1.00 19.75
                                   47.533
                                            -9.507
                   VAL B 175
ATOM
        4392
              CG1
                                            -7.169
                                                     81.396
                                                              1.00 18.66
                   VAL B 175
                                   46.896
        4393
              CG<sub>2</sub>
MOTA
                                                              1.00 22.82
                                   46.635 -10.486
                                                     83.324
                   VAL B 175
MOTA
        4394
              C
                                   46.255 -11.503
                                                     82.754
                                                              1.00 18.06
                   VAL B 175
        4395
              O
MOTA
                                   47.439 -10.520
                                                     84.378
                                                              1.00 21.67
        4396
              N
                   GLN B 176
MOTA
                                                              1.00 21.55
                   GLN B 176
                                   47.889 -11.798
                                                     84.911
        4397
              CA
ATOM
                                                     86,105
                                                              1.00 19.68
                   GLN B 176
                                   48.824 -11.602
        4398
              CB
ATOM
                                   49.088 -12.905
                                                     86.862
                                                              1.00 20.17
MOTA
        4399
                   GLN B 176
              CG
                                                     87.996
                                                              1.00 25.42
                                   50.066 -12.759
                   GLN B 176
        4400
              CD
MOTA
                                                     87.786
                                   51.243 -12.442
                                                              1.00 21.56
        4401
              OE1
                   GLN B 176
ATOM
                                                              1.00 20.18
                                   49.592 -13.000
                                                     89.217
                   GLN B 176
        4402
              NE<sub>2</sub>
ATOM
                                                     85.348
                                                              1.00 24.78
                                   46.689 -12.630
                   GLN B 176
        4403
              С
ATOM
                                                              1.00 22.91
                                                     85.057
                                   46.618 -13.817
              0
                   GLN B 176
        4404
MOTA
                                   45.751 -12.007
                   GLU B 177
                                                     86.051
                                                              1.00 23.69
        4405
              N
ATOM
                                   44.571 -12.727
                                                     86.523
                                                              1.00 27.01
                   GLU B 177
        4406
ATOM
              CA
                                                              1.00 24.73
                                                     87.394
                                   43.703 -11.825
                   GLU B 177
        4407
              CB
MOTA
                   GLU B 177
                                   42.633 -12.581
                                                     88.138
                                                              1.00 37.46
              CG
        4408
MOTA
                                   41.767 -11.676
                                                     88.987
                                                              1.00 42.48
                   GLU B 177
        4409
              CD
ATOM
                                                     88.432
                                                              1.00 44.35
                                   40.875 -11.002
                   GLU B 177
        4410
              OE1
ATOM
                                                              1.00 45.63
              OE2
                   GLU B 177
                                   41.993 -11.627
                                                     90.213
        4411
MOTA
                   GLU B 177
GLU B 177
                                   43.732 -13.247.
                                                     85.370
                                                              1.00 26.56
        4412
              C
MOTA
                                   43.240 -14.375
                                                              1.00 27.71
                                                     85.408
ATOM
        4413
              0
                                                     84.344
                                                              1.00 24.58
        4414
              N
                   ALA B 178
                                   43.573 -12.418
ATOM
                                                              1.00 25.86
                                                     83.174
                   ALA B 178
                                   42.776 -12.775
        4415
              CA
MOTA
                   ALA B 178
                                   42.778 -11.628
                                                     82.171
                                                              1.00 24.20
MOTA
        4416
              CB
                                   43.231 -14.054
                                                                   25.72
                                                     82.485
                                                              1.00
        4417
                   ALA B 178
ATOM
                                                              1.00 22.38
                                                     82.036
                   ALA B 178
                                   42.406 -14.838
        4418
              0
MOTA
                                                              1.00 27.19
                   PHE B 179
                                   44.535 -14.282
                                                     82.395
              N
ATOM
        4419
                                                     81.703
                                                              1.00 27.05
                   PHE B 179
                                   44.990 ~15.489
        4420
              CA
MOTA
                                                              1.00 25.22
                                   45.714 -15.086
                                                     80.418
              CB
                   PHE B 179
        4421
MOTA
                                                              1.00 20.36
                                                     79.644
                   PHE B 179
                                   44.992 -14.020
              CG
        4422
MOTA
                                                     79.735
                                                              1.00 25.23
                  PHE B 179
                                   45.387 -12.687
        4423
              CD1
MOTA
                                   43.860 -14.332
                                                     78.902
                                                              1.00 19.22
                  PHE B 179
        4424
              CD2
ATOM
                                                     79.102
                                                              1.00 19.25
                                   44.659 -11.677
                  PHE B 179
              CE1
        4425
ATOM
                                                              1.00 20.65
                                   43.128 -13.315
                                                     78.272
              CE2
                  PHE B 179
        4426
ATOM
                                                     78.374
                                                              1.00 25.64
                                   43.528 -12.001
                  PHE B 179
        4427
              cz
MOTA
                                                     82.556
                                                              1.00 23.50
                                   45.866 -16.398
                   PHE B 179
        4428
              С
MOTA
                                                              1.00 18.26
                                                     82.038
                                   46.652 -17.182
                   PHE B 179
        4429
              0
ATOM
                                   45.689 -16.313
                                                     83.868
                                                              1.00 23.24
                   TYR B 180
        4430
              N
MOTA
                                                     84.799
                                                              1.00 26.76
                                   46.479 -17.106
                  TYR B 180
        4431
              CA
MOTA
                                                     86.231
                                                              1.00 25.72
                  TYR B 180
                                   46.150 -16.665
              CB
       4432
MOTA
                                                              1.00 29.66
                                                     87.247
                  TYR B 180
                                   47.226 -16.969
        4433
              CG
MOTA
                                   47.037 -17.942
                                                     88.237
                                                              1.00 27.07
                  TYR B 180
        4434
              CD1
MOTA
                                                     89.170
                                                              1.00 30.08
                  TYR B 180
                                   48.039 -18.222
              CE1
       4435
MOTA
                                                              1.00 29.68
                                                     87.216
                  TYR B 180
                                   48.444 -16.283
       4436
              CD2
MOTA
                                   49.451 -16.552
                                                     88.139
                                                              1.00 30.99
                  TYR B 180
       4437
              CE2
ATCM
                                                     89.112
                                                              1.00 33.16
                  TYR B
                        180
                                   49.248 -17.521
              CZ
       4438
ATOM
                                                              1.00 28.47
                                                     90.006
                                   50.262 -17.791
                  TYR B 180
       1439
              OH
ATOM
                                   46.256 -18.619
                                                     84.649
                                                              1.00 29.13
                  TYR B 180
       4440
MOTA
                                                     84.922
                                                              1.00 23.43
                        180
                                   47.163 -19.416
                  TYR B
       4441
              0
MOTA
                                                              1.00 25.67
                                                     84.190
                                   45.073 -19.021
                  ASP B 181
       4442
              N
ATOM
                                   44.784 -20.445
                                                     84.075
                                                              1.00 28.28
                  ASP B 181
              CA
       4443
MOTA
                                                     84.757
                                                              1.00 32.13
                  ASP B
                                   43.446 -20.759
                         181
              CB
       1444
ATOM
                                                              1.00 36.12
                                                     83.890
                                   42.247 -20.410
                  ASP B 181
              CG
       4445
ATCM
```

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								•
MOTA	4446	OD1	ASP B 1	81	42.20	2 -19.300	83.329	1.00 41.04
ATOM	4447			81		4 -21.249		
ATOM	4448		ASP B 1			3 -21.018		
	4449		ASP B 1					
MOTA						6 -22.115		
ATOM	4450		THR B 1			5 -20.302		
ATOM	4451		THR B 1			3 -20.823		1.00 30.57
MOTA	4452	CB _ 1	THR B 1	32	44.46	8 -20.008	79.397	1.00 30.03
MOTA	4453	OG1	THR B 1	32	44.51	6 -20.598	78.095	1.00 28.22
MOTA	4454	CG2 5	THR B 1	32	44.94	7 -18.561	79.310	
MOTA	4455	C :	THR B 1	32		9 -20.870		
ATOM	4456		THR B 1			1 -20.007	80.008	
ATOM	4457		ASP B 1			9 -21.878	78.909	1.00 27.27
MOTA	4458		ASP B 1			6 -22.049		
							78.273	1.00 31.40
MOTA	4459		ASP B 1			8 -23.536	78.228	1.00 33.36
ATOM	4460		SP B 1			8 -24.319	77.328	1.00 33.33
ATOM	4461		ASP B 1			3 -23.988	77.28 7	1.00 28.06
ATOM	4462		SP B 18			5 -25.271	76.675	1.00 38.19
ATOM	4463		ASP B 18			L -21.462	76.864	1.00 31.14
MOTA	4464	O A	SP B 18	3	49.332	2 -21.557	76.168	1.00 28.74
MOTA	4465	N G	LN B 18	4	47.217	7 -20.852	76.446	1.00 25.34
ATOM	4466	CA G	LN B 18	4	47.151	-20.251	75.118	1.00 28.59
MOTA	4467	CB G	LN B 18	4		-20.256	74.581	1.00 26.84
ATOM	4468		LN B 18			-21.632	74.529	1.00 34.86
ATOM	4469		LN B 18			-21.647	73.736	1.00 32.27
MOTA	4470		LN B 18			-20.789	73.730	1.00 35.43
ATOM	4471		LN B 18			-22.641	72.870	
ATOM	4472		LN B 18			-18.817		1.00 28.92
	4473						75.175	1.00 27.28
MOTA						-18.171	74.148	1.00 29.70
ATOM	4474		AL B 18			-18.325	76.386	1.00 27.64
MOTA	4475		AL B 18	-		-16.972	76.575	1.00 26.26
MOTA	4476		AL·B 18			-16.039	77.145	1.00 22.85
MOTA	4477	CG1 V				-14.642	77.395	1.00 23.10
MOTA	4478	CG2 V		_		-15.967	76.191	1.00 21.67
ATOM	4479		AL B 18			-16.964	77.547	1.00 27.01
ATOM	4480		AL B 18		49.456	-17.469	78.663	1.00 23.75
ATOM	4481	N P	HE B 18	5.	50.696	-16.403	77.115	1.00 22.02
ATOM	4482	CA PI	HE B 18	5	51.868	-16.301	77.978	1.00 21.83
MOTA	4483	CB PI	HE B 18	5	53.142	-16.763	77.252	1.00 17.02
ATOM	4484	CG PI	HE B 18	5	54.336	-16.921	78.170	1.00 24.84
ATOM	4485	CD1 PI	HE B 18	5	54.756	-18.189	78.580	1.00 22.70
ATOM	4486	CD2 PI	HE B 18	5	55.004	-15.805	78.670	1.00 20.26
ATOM	4487	CE1 PH	E B 18	5		-18.338	79.471	1.00 21.47
ATOM	4488		IE B 18			-15.941	79.563	1.00 20.01
ATOM	4489		E B 18			-17.206	79.968	1.00 17.84
ATOM	4490		E B 18			-14.827	78.368	1.00 18.12
ATO:	4491		E B 18			-13.946	77.508	1.00 15.12
ATO.4	4492		T B 18.				79.661	
	4493		L B 187			-14.565		1.00 18.06
ATON						-13.208	80.153	1.00 17.67
ATOM	4494		L B 187			-12.839	81.225	1.00 22.85
MOTA	4495		L B 187			-11.473	81.840	1.00 24.08
MOTA	4496		L B 187			-12.808		
MOTA	4497		L B 187			-13.060	80.788	1.00 18.32
MOTA	1498	O VA	L B 187			-13.807	81.707	1.00 18.82
ATOM	4499		T B 188		54.503	-12.103°	80.282	1.00 14.70
MOTA	4500	CA LE	U B 188			-11.789	80.798	1.00 18.84
ATCM	4501		U B 188		56.900	-11.948	79.716	1.00 18.64
MOTA	4502	CG LE	U B 188			-11.277	80.082	1.00 21.23
ATOM	4503		U B 188			-11.832	81.395	1.00 18.55
ATOM	4504		U B 188			-11.489	78.957	1.00 20.49
ATOM	4505		U B 188			-10.339	81.280	1.00 22.14
ATOM	4506		U B 188		55.527	-9.410	80.517	1.00 19.96
ATOM	4507		R B 189			-10.133	82.540	1.00 21.08
	4508		R B 189		56.203	-8.782	83.061	1.00 21.05
ATOM	4509		R B 189		54.956		83.908	1.00 25.95
MOTA					54.988	-8.543		
ATCM	4510		R B 189			-7.252	84.475	1.00 21.91
ATOM.	4511	C SE	R B 189		57.423	-8.420	83.883	1.00 23.62

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ATOM	4512	0	.SER B	189		57.829	-9.174	84.766	1.00 18.61
ATOM	4513	N	LEU B	190		58.020	-7.269	83.569	1.00 20.83
	4514	CA	LEU B			59.149	-6.767	84.347	1.00 21.85
ATOM			LEU B			60.278	-6.226	83.473	1.00 22.85
MOTA	4515	CB							1.00 32.59
ATOM	4516	CG	LEU B			60.964	-7.089	82.413	
MOTA	4517	CD1	LEU B	190		62.337	-6.479	82.140	1.00 29.27
ATOM	4518	CD2	LEU B	190		61.136	-8.511	82.879	1.00 31.88
MOTA	4519	С	LEU B	190		58.505	-5.613	85.085	1.00 21.28
	4520	ō	LEU B			57.695	-4.897	84.501	1.00 15.72
MOTA			HIS B			58.857	-5.421	86.351	1.00 18.16
ATOM	4521					58.249	-4.357	87.145	1.00 17:46
MOTA	4522	CA	HIS B						1.00 17.40
МОТА	4523	CB	HIS B	-		56.759	-4.690	87.369	
MOTA	4524	CG	HIS B	191	-	56.517	-6.085	87.880	1.00 22.14
ATOM	4525	CD2	HIS B	191		56.341	-6.551	89.143	1.00 12.25
ATOM	4526	ND1	HIS B	191		56.372	-7.179	87.049	1.00 18.02
ATOM	4527		HIS B			56.119	-8.256	87.775	1.00 8.17
	4528	NE2				56.094	-7.902	89.049	1.00 19.79
ATOM			HIS B			58.945	-4.197	88.484	1.00 17.41
ATOM	4529	С					-5.029	88.867	1.00 18.74
MOTA	4530	0	HIS B			59.769			
MOTA:	4531	N	GLN B			58.618	-3.114	89.182	1.00 18.20
MOTA	4532	CA	GLN B	192		59.173	-2.854	90.502	1.00 18.41
ATOM	4533	CB	GLN B	192		58.690	-1.500	91.034	1.00 20.71
ATOM	4534	CG	GLN B	192		58.871	-0.334	90.072	1.00 21.49
ATOM	4535	CD	GLN B			58.226	0.930	90.594	1.00 20.65
	4536	OE1				58.775	1.615	91.459	1.00 21.52
MOTA		NE2				57.029	1.226	90.098	1.00 15.10
ATOM	4537						-3.945	91.395	1.00 17.55
ATOM	4538	С	GLN B	192		58.608		91.320	1.00 17.33
ATOM	4539	0	GLN B	192	•	57.415	-4.256		
ATOM	4540	N	SER B			59.447	-4.522	92.240	1.00 15.71
ATOM	4541	CA	SER B	193		58.986	-5.574	93.143	1.00 20.58
ATOM	4542	CB	SER B	193		60.093	-5.963	94.120	1.00 20.71
	4543	OG	SER B	193		59.571	-6.804	95.138	1.00 22.55
MOTA	4544	c	SER B	193		57,774	-5.112	93.947	1.00 21.81
ATOM			SER B	193		57.769	-4:003	94.486	1.00 20.82
ATOM ·	4545	0				56.745	-5.967	94.063	1.00 21.80
MOTA	4546	N	PRO B	194				93.524	1.00 24.27
MOTA	4547	CD	PRO B			56.648	-7.331		
MOTA	4548	CA	PRO B			55.524	-5.643	94.812	1.00 23.58
ATOM	4549	CB	PRO B	194		54.678	-6.909	94.642	1.00 22.98
ATOM	4550	CG	PRO B	194		55.168	-7.458	93.317	1.00 26.35
ATOM .	4551	С	PRO B	194		55.841	-5.366	96.283	1.00 25.79
	4552	0	PRO B	194		55.009	-4.831	97.022	1.00 27.26
ATOM	4553	N	GLU B	195		57.045	-5.736	96.710	1.00 23.20
MOTA			GLU B			57.428	-5.514	98.093	1.00 29.56
MOTA	4554	CA					-6.090	98.379	1.00 32.38
ATOM	4555	CB	GLU B	195		58.816			1.00 45.25
ATOM	4556	CG	GLU B	195		58.940	-7.567		
ATOM	4557	CD	GLU B	195		60.206	-8.189	98.613	1.00 50.44
MOTA	4558	OE1	GLU B	195		61.290	-7.580	98.471	1.00 50.51
ATOM	4559	OE2	GLU B	195		60.118	-9.297	99.184	.00 49.77
ATOM	4560	С	GLU B	195		57.414	-4.035	98.425	1.00 25.11
	4561	ō				57.095	-3.659	99.551	1.00 29.05
ATOM	4562	N	TYR B			57.729	-3.191	97.445	1.00 22.90
ATOM						57.743	-1.750	97.696	1.00 22.46
ATOM	4563	CA	TYR B		•		-1.223	97.668	1.00 22.72
ATCM	4564	CB	TYR B			59.188			1.00 24.17
ATOM	.4565	CG	TYR B	196		59.855	-1.234	96.301	
ATOM	4566	CD1	TYR B			59.639	-0.203	95.385	1.00 20.87
ATOM	4567	CE1	TYR B	196		60.229	-0.222	94.118	1.00 18.31
	4568	CD2	TYR B			60.684	-2.289	95.916	1.00 24.63
ATOM	4569	CE2	TYR B			61.276	-2.318	94.648	1.00 24.39
ATOM		CZ	TYR B			61.042	-1.284	93.756	1.00 23.01
MOTA	4570					61.592	-1.328	92.492	1.00 19.86
ATOM	4571	ОН	TYR B					96.725	1.00 23.54
ATOM	4572	С	TYR B			56.896	-0.938		T.00 73.74
ATOM	4573	0	TYR B			56.779	0.275	96.869	1.00 17.53
ATOM	4574	N	ALA B	197		56.293	-1.589	95.740	1.00 22.11
ATOM	4575	CA		197		55.503	-0.829	94.779	1.00 24.28
	4576	CB		197		56.310	-0.616	93.513	1.00 23.03
ATOM	1577	C	ALA B			54.153	-1.412	94.413	1.00 22.80
ATOM	45//	_	מ אונה		_	J-2 - #		-	

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Figure 18-70

MOTA	4578	0	ALA B 197	53.910	-2.609	94.549	1.00 17.67
MOTA	4579	N	PHE B 198	53.278		93.932	1.00 26,40
ATOM	4580	CA	PHE B 198	51.956	-0.950	93.495	1.00 28,19
ATOM	4581			51.152		93.035	1.00 29.51
MOTA	4582			49.721		92.711	1.00 29.50
ATOM	4583			48.732	0.1.00	93.674	1.00 32.77
ATOM	4584			49.367		91.455	1.00 25.82
MOTA	4585			47.410		93.394	1.00 36.70
MOTA	4586			48.050		91.170	1.00 29.29
	4587			47.071	-0.703	92.141	1.00 23.23
MOTA	4588		PHE B 198	52.170	-1.858	92.284	1.00 28.28
MOTA	4589		PHE B 198	53.045	-1.602	91.456	1.00 27.15
ATOM			PRO B 199			92.185	
ATOM	4590			51.407	-2.952		1.00 31.37
MOTA	4591			51.440	-3.887	91.045	1.00 37.07
MOTA	4592		PRO B 199	50.386	-3.369	93.144	1.00 35.32
MOTA	4593	CB	PRO B 199	49.545	-4.328	92.321	1.00 33.88
MOTA	4594	CG	PRO B 199	50.641	-5.068	91.578	1.00 36.75
MOTA	4595	C	PRO B 199	51.241	-4.082	94.184	1.00 36.93
MOTA	4596	0	PRO B 199	52.308	-4.603	93.860	1.00 50.93
MOTA	4597	N	PHE B 200	50.804	-4.127	95.422	1.00 37.04
MOTA	4598	CA	PHE B 200	51.644	-4.763	96.421	1.00 30.13
ATOM	4599	CB	PHE B 200	51.547	-3.968	97.723	1.00 28.70
MOTA	4600	CG	PHE B 200	51.760	-2.485	97.543	1.00 29.98
MOTA	4601		L PHE B 200	50.717	-1.660	97.137	1.00 28.92
ATOM	4602		2 PHE B 200	53.016	-1.919	97.746	1.00 23.60
MOTA	4603	CE:		50.922	-0.289	96.938	1.00 27.63
MOTA	4604	CEZ		53.229	-0.558	97.547	1.00 23.56
MOTA	4605	CZ	PHE B 200	52.182	0.260	97.143	1.00 28.37
ATOM	4606	C	PHE B 200	51.296	-6.227	96.658	1.00 25.51
ATOM	4607	0	PHE B 200	52.112	-6.984	97.167	1.00 20.92
ATOM	4608	N	GLU B 201	50.094	-6.618	96.252	1.00 27.41
ATOM	4609	CA	GLU B 201	49.576	-7.972	96.454	1.00 31.98
MOTA	4610	CB	GLU B 201	48.056	-7.928	96.487	1.00 31.57
MOTA	4611	CG	GLU B 201	47.486	-6:935	97.449	1.00 39.17
MOTA	4612	CD	GLU B 201	45.987	-6.853	97.316	1.00 40.31
ATOM	4613	OE1	GLU B 201	45.332	-7.902	97.500	1.00 38.90
MOTA	4614	OE2	GLU B 201	45.475	-5.751	97.019	1.00 35.04
MOTA	4615	С	GLU B 201	49.979	-9.018	95.422	1.00 30.83
MOTA	4616	0	GLU B 201	49.901	-10.219	95.690	1.00 26.34
MOTA	4617	N	LYS B 202	50.362	-8.573	94.234	1.00 24.95
MOTA	4618	CA	LYS B 202	50.764	-9.501	93.195	1.00 22.79
ATOM	4619	CB	LYS B 202	49.588	-9.773	92.258	1.00 25.12
MOTA	4620	CG	LYS B 202	48.484	-10.523	93.000	1.00 35.38
MOTA	4621	CD	LYS B 202	47.431	-11.099	92.103	1.00 38.67
MOTA	4622	CE	LYS B 202	46.498	-11.998	92.903	1.00 40.98
ATOM	4623	NZ	LYS B 202	.45.491	-12.659	92.028	1.00 46.65
MOTA	4624	С	LYS B 202	51.975	-9.007	92.435	1.00 24:62
MOTA	4625	0	LYS B 202	52.355	-7.838	92.549	1.00 21.83
MOTA	4626	N	GLY B 203	52.598	-9.910	91.684	1.00 17.60
ATOM	4627	CA	GLY B 203	53.779	-9.545	90.928	1.00 19.41
ATOM	4628	C	GLY B 203		-10.297	91.396	1.00 20.36
ATOM	4629	Ō	GLY B 203		-10.070	90.888	1.00 23.83
ATOM	4630	N	PHE B 204		-11.201	92.358	1.00 24.82
MOTA	4631	CA	PHE B 204		-11.957	92.859	1.00 24.24
ATOM	4632	CB	PHE B 204		-12.567	94.236	1.00 22.72
ATOM	4633	CG	PHE B 204	55.485		95.322	1.00 25.26
ATOM	4634		PHE B 204	54.235		95.535	1.00 23.80
	4635			56.551		96.133	1.00 20.25
MOTA		CE1	PHE B 204		-10.036	96.545	1.00 20.23
MOTA	4636		PHE B 204	56.377		97.141	1.00 24.40
ATOM	4637	CE2	PHE B 204			97.350	1.00 25.54
ATOM	4638	CS		55.124	-9.658	91.894	1.00 25.86
MOTA	4639	C	PHE B 204	56.412		91.091	
ATOM	4640	0	PHE B 204	55.613		91'.986	1.00 20.65
MOTA	1641	N	LEU B 205	57.676			1.00 24.25
ATOM	4642	CA	LEU B 205	58.233		91.114	1.00 30.66
ATOM	4643	CB	LEU B 205	59.723	-14.637	91.413	1.00 34.01

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Figure 18-71

```
60.495 -15.669
                                                    90,592
                                                             1.00 34.12
MOTA
        4644
               CG
                   LEU B 205
                   LEU 3 205
                                   60.356 -15.382
                                                     89.109
                                                             1.00 32.95
ATOM
        4645
               CD1
                   LEU B 205
                                   61.957 -15.629
                                                     91,005
                                                             1.00 36.49
MOTA
        4646
               CD2
MOTA
        4647
                   LEU B 205
                                   57.535 -15.827
                                                     91.205
                                                             1.00 30.51
                                   57.467 -16.562
        4648
               0
                   LEU B 205
                                                     90.220
                                                             1.00 25.89
ATOM
                                   57.010 -16.147
                                                     92.382
                                                             1.00 30.43
                   GLU B 206
ATOM
        4649
               N
                   GLU B 206
                                   56.338 -17.423
                                                     92.605
                                                             1.00 30.64
ATOM
        4650
               CA
        4651
               CB
                   GLU B 206
                                   56.025 -17.601
                                                     94.093
                                                             1.00 34.77
MOTA
                   GLU B 206
                                                             1.00 42.50
                                   57.227 -17.512
                                                     95.033
MOTA
        4652
               CG
        4653
                   GLU B 206
                                   57.718 -16.084
                                                     95.270
                                                             1.00 45.76
ATOM
               CD
        4654
               OE1
                   GLU B 206
                                   58.228 -15.438
                                                     94.333
                                                             1.00 42.62
ATOM
                                   57.585 -15.602
                                                             1.00 50.22
                   GLU B 206
                                                     96,413
MOTA
        4655
               OE2
                   GLU B 206
                                   55.045 -17.587
                                                     91.811
                                                             1.00 31.13
ATOM
        4656
               C
        4657
                   GLU B 206
                                   54.607 -18.708
                                                     91.563
                                                             1.00 28.18
MOTA
              Ω
                                                     91.425
                                                             1.00 25.16
                   GLU B 207
                                   54.430 -16.472
MOTA
        4658
              N
                                                             1.00 28.78
                   GLU B 207
                                   53.178 -16.499
                                                     90.664
MOTA
        4659
               CA
                   GLU B 207
                                   52.546 -15.107
                                                     90.695
                                                             1.00 30.76
        4660
              CB
ATOM
                                   52.121 -14.659
                                                     92.093
                                                             1.00 29.39
                   GLU B 207
ATOM
        4661
              CG
                                   52.057 -13.151
                                                             1.00 27.87
MOTA
        4662
              CD
                   GLU B 207
                                                     92.230
        4663
              OE1 GLU B 207
                                   51.656 -12.477
                                                     91.261
                                                             1.00 24.38
ATOM
                  GLU B 207
                                   52.389 -12.636
                                                     93.316
                                                             1.00 25.36
MOTA
        4664
              OE2
                                                             1.00 29.48
                                   53.453 -16.922
MOTA
        4665
                   GLU B 207
                                                     89.224
        4666
              0
                   GLU B 207
                                   53.658 -16.077
                                                     88.351
                                                             1.00 27.48
MOTA
                   ILE B 208
                                   53.442 -18.230
                                                    88.976
                                                             1.00 26.67
        4667
MOTA
              N
                                                             1.00 32.60
                                   53.735 -18.754
                                                     87.646
MOTA
        4668
              CA
                   ILE B 208
ATOM
        4669
              CB
                   ILE B 208
                                   54.789 -19.877
                                                    87.740
                                                             1.00 34.26
                                   55.239 -20.296
                                                    86.352
                                                             1.00 41.65
              CG2
                   ILE B 208
        4670
ATOM
                                   56.008 -19.404
                         208
                                                    88.532
                                                             1.00 36.07
        4671
                   ILE B
MOTA
              CG1
                                                             1.00 45.18
        4672
              CD1 ILE B 208
                                   56.814 -18.338
                                                    87.851
ATOM
                         208
                                   52.522 -19.289
                                                    86.870
                                                             1.00 32.26
        4673
              С
                   ILE B
MOTA
                                                    85.759
                                                             1.00 27.43
        4674
                   ILE B 208
                                   52.668 -19.799
              0
MOTA
                                   51.328 -19.165
                                                    87.442
                                                             1.00 32.60
        4675
ATOM
              N
                   GLY B 209
        4676
              CA
                   GLY B 209
                                   50.139 -19.652
                                                    86.760
                                                             1.00 35.07
MOTA
                   GLY B 209
                                   49.565 -20.892
                                                    87.420
                                                             1.00 36.19
        4677
ATOM
                   GLY B 209
                                   50.230 -21.524
                                                    88.235
                                                             1.00 31.61
        4678
ATOM
              0
                                                             1.00 36.98
                  GLU B 210
                                   48.335 -21.245
                                                    87.066
ATOM
        4679
              N
                                                             1.00 40.60
ATOM
        4680
                   GLU B 210
                                   47.677 -22.412
                                                    87.647
              CA
                                   46.633 -21.964
                                                             1.00 37.98
        4681
              CB
                   GLU B 210
                                                    88.672
MOTA
                  GLU B 210
                                   45.446 -21.234
                                                    88.058
                                                             1.00 42.78
ATOM
        4682
              CG
                                                             1.00 48.41
        4683
              CD
                   GLU B 210
                                   44.470 -20.717
                                                    89.098
ATOM
                      B 210
                                   43.400 -20.202
                                                    88.709
                                                             1.00 51.03
        4684
              OE1
                  GLU
ATOM
                  GLU B 210
                                  44.778 -20.814
                                                    90.306
                                                             1.00 49.90
        4685
              OE2
ATOM
                   GLU B 210
                                  46.996 -23.248
                                                    86.564
                                                             1.00 39.48
        4686
              С
ATOM
                  GLU B 210
                                   46.709 -22.751
                                                    85.471
                                                             1.00 33.65
        4687
ATOM
                                   46.736 -24.515
                                                    86.876
                                                             1.00 39.18
        4.688
              N
                  GLY B 211
ATOM
                  GLY B 211
                                  46.087 -25.399
                                                    85.923
                                                             1.00 38.43
        4689
MOTA
              CA
                                                             1.00 40.29
        4690
                  GLY B 211
                                   46,877 -25.500
                                                    84.637
MOTA
                  GLY B 211
                                   48.101 -25.610
                                                    84.666
                                                             1.00 39.39
        4691
MOTA
        4692
                                  46.187 -25.458
                                                    83.504
                                                             1.00 40.90
              N
                  LYS B 212
MOTA
                                                             1.00 43.53
                                  46.864 -25.538
MOTA
        4693
              CA
                  LYS B 212
                                                    82.219
                  LYS B 212
                                                             1.00 47.87
                                  45.842 -25.548
                                                    81.080
ATOM
        4694
              CB
       4695
              CG
                  LYS B 212
                                  44.795 -26.665
                                                    81.144
                                                             1.00 53.09
ATOM
                                  45.398 -28.076
                                                             1.00 58.61
                                                    81.130
ATOM
        4696
              CD
                  LYS B 212
                                                             1.00 59.78
       4697
                  LYS B 212
                                  46.069 -28.454
                                                    82.452
ATOM
              CE
       4698
                                  46.670 -29.825
                                                    82.420
                                                             1.00 62.17
                  LYS B 212
ATOM
              NZ
                                  47.823 -24.363
                                                    82.040
                                                             1.00 38.84
                  LYS B 212
MOTA
       4699
                                                             1.00 40.33
ATOM
       4700
                  LYS B 212
                                  48.797 -24.457
                                                    81.295
                  GLY B 213
GLY B 213
                                  47.543 -23.262
                                                    82.731
                                                             1.00 37.20
       4701
              N
ATOM
                                                    82.627
                                                             1.00 34.66
                                  48.384 -22.081
ATOM
       4702
              CA
                                                             1.00 37.09
       4703
                  GLY B 213
                                  49.625 -22.107
                                                    83.505
ATOM
              С
       4704
                                  50.425 -21.165
                                                    83.489
                                                             1.00 25.85
                  GLY B 213
ATOM
              0
                                                    84.273
                                                             1.00 33.33
                                  49.794 -23.180
       4705
              N
                  LY5
                      в 214
ATOM
                                                             1.00 37.90
       4706
                  LYS B 214
                                  50.953 -23.297
                                                    85.148
              CA
MOTA
       4707
                  LYS B 214
                                  50.886 -24.598
                                                    85.954
                                                             1.00 38.89
              CB
ATCM
                                                             1.00 39.29
                  LYS B 214
                                                    86.938
                                  52.032 -24.786
ATOM
       4708
              CG
                                                            1.00 43.60
                                                    87.704
       4709
                  LYS B 214
                                  51.876 -26.094
ATCM
```

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Figure 18-72

	4710	CE	7 37 6	B 214		E2 047	-26.334	88.640	1 00	47.36
atom	4710	CE								
MOTA	4711	NZ		B 214		53.165		89.666		54.03
MOTA	4712	С	LYS	B 214	l.	52.209	-23.275	84.291	1.00	34.16
MOTA	4713	0	LYS	B 214	l	52.404	-24.136	83.438	1.00	34.70
			GLY			53.057		84.523		33.58
MOTA	4714	N								
MOTA	4715	CA	GLY			54.275		83.743		28.27
MOTA	4716	С	GLY	B 215	;	54.104	-21.155	82.605	1.00	31.02
ATOM	4717	ō	GLY			55.033	-20.911	81.833	1 00	23.68
-										22.45
ATOM	4718	N	TYR			52.918		82.493		
MOTA	4719	CA	TYR	B 216)	52.683	-19.605	81.426	1.00	24.03
MOTA	4720	CB	TYR	B 216	· .	51.458	-20.013	80.603	1.00	17.60
	4721	CG	TYR			51.682		79.806		25.96
MOTA										
MOTA	4722	CD1					-22.538	80.435		21.41
MOTA	4723	CE1	TYR	B 216		51.988	-23.704	79.729	1.00	23.78
ATOM	4724	CD2	TYR	B 216	;	51.970	-21,242	78.439	1.00	19.62
							-22.402	77.72-2		26.39
MOTA	4725	CE2								
MOTA	4726	CZ	TYR	B 216	٠.		-23.630	78.379		29.35
ATOM	4727	OH	TYR	B 216		52.577	-24.782	77.690	1.00	27.75
ATOM	4728	C		B 216			-18.153	81.884	1.00	24.53
				B 216			-17.298	81.159		18.99
ATOM	4729	0								
MOTA	4730	N	ASN	B 217		53.052	-17.886	83.098		21.41
MOTA	4731	CA	ASN	B 217		53.073	-16.534	83.642	1.00	21.23
ATOM	4732	СВ		B 217			-16.325	84.669	1.00	16.78
MOTA	4733	CG		B. 217			-14.889	85.162		22.07
MOTA	4734	OD1	ASN	B 217		52.506	-14.521	86.163	1.00	23.13
ATOM	4735	ND2	ASN	B 217		51.146	-14.058	84.435	1.00	19.26
ATOM	4736	С		B 217			-16.339	84.291	1.00	19.40
								85.124	1.00	
MOTA	4737	0		B 217			-17.145			
MOTA	4738	N	LEU	B 218			-15.273	83.905		18.65
MOTA	4739	CA	LEU	B 218		56.459	-15.004	84.444	1.00	16.41
ATOM	4740	CB	LEU			57.512	-15.244	83.368	1.00	18.29
							-15.872	83.782		28.15
MOTA	4741	CG	LEU							
ATOM	4742		LEU				-15.563	82.695		20.50
MOTA	4743	CD2	LEU	B 218		59.332	-15.348	.85.116	1.00	22.53
MOTA	4744	Ċ	LEU			56.595	-13.562	84.926	1.00	17.89
	4745	ō	LEU				-12.627	84.128		14.48
MOTA										
MOTA	4746	N	ASN				-13.395	86.219		14.09
ATOM	4747	CA	ASN	B 219		57.044	-12.075	86.821	1.00	18.41
ATOM	4748	CB	ASN	B 219		56.238	-11.922	88.111	1.00	14.64
	4749	CG	ASN				-11.898	87.868		27.12
ATOM										
MOTA	4750		ASN				-11.332	86.880		20.21
ATOM	4751	ND2	ASN	B 219		53.982	-12.480	88.787	1.00	23.62
MOTA	4752	С	ASN .	B 219		58.504	-11.843	87.172	1.00	20.39
	4753	ō	ASN				-12.672	87.841	1.00	20.41
MOTA										15.11
MOTA	4754	N		B 220			-10.717	86.729		
MOTA	4755	CA	ILE :	B22C		60.441	-10.394	87.033		17.16
ATOM	4756	CB	ILE :	B 220		61.250	-10.083	85.740	1.00	20.78
	4757	CG2				62.736	-9.821	86.094	1.00	18.08
MOTA								84.748		17.62
MOTA	4758	CG1	ILE :				-11.250			
ATOM	4759	CD1	ILE :	B 220		61.646	-12.590	85.273		20.72
ATOM	4760	С	ILE 1	B 220		60.475	-9.161	87.947		21.17
	4761	Ó		B 220		60.565	-8.036	87.470	.1.00	16.03
MOTA								89.274		21.74
ATOM	4762	N		B 221		60.367	-9.357			
MOTA	4763	CD	PRO 1	B 221		60.135	-10.619	90.000		22.96
ATOM	4764	CA	PRO I	B 221		60.394	-8.225	90.213	1.00	19.16
	4765	CB		3 221		59.947	-8.869	91.523		19.40
MOTA										23.02
ATOM	4766	CG		B 221		60.564	-10.251	91.407		
MOTA	4767	С	PRO I	3 221		61.799	-7.634	90.289	1.00	22.42
MOTA	4768	0	PRO 1	3 221		62.780	-8.369	90.425	1.00	20.71
		N		3 222		61.899	-6.309	90.202		22.74
ATOM	4769									
ATOM	4770	CA		3 222		63.198	-5.643	90.223	1.00	21.18
ATOM	4771	CB	LEU I	3 222		63.453	-4.993	88.850	1.00	17.21
ATOM	4772	CG		3 222		63.467	-6.027	87.721	1.00	20.26
			LEU I			63.453	-5.354	86.361	1 00	20.00
ATCM	4773								1 00	21.93
ATCM	4774		LEU I			64.696	-6.908	.87.881	1.00	21.73
ATOM	4775	С	LEU I	3 222		63.335	-4.616	91.353	1.00	20.04
								-		

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Figure 18-73

```
91.806
                                                              1.00 17.58
                                            -4.030
ATOM
        4776
               0
                   LEU B 222
                                    62.350
                                                             . 1.00 19.48
                   PRO B 223
                                    64.571
                                            -4.394
                                                     91.830
        4777
ATOM
               N
                   PRO B 223
                                    65.806
                                            -5.072
                                                     91.400
                                                              1.00
                                                                   16.80
MOTA
        4778
               CD
                                                     92.915
                                                              1.00 20.38
                                    64.873
MOTA
        4779
               CA
                   PRO B 223
                                            -3.454
                                                              1.00 26.11
                   PRO B 223
                                    66.274
                                            -3.881
                                                     93.327
               CB
        4780
MOTA
                   PRO B 223
                                                     91.973
                                                              1.00 19.74
                                    66.884
                                            -4.161
MOTA
        4781
               CG
                                                     92.553
                                            -1.971
                                                              1.00 21.39
                   PRO B 223
                                    64.818
        4782
               С
MOTA
                                                              1.00 17.16
                   PRO B 223
                                    64.815
                                            -1.598
                                                     91.380
        4783
               O
MOTA
                   LYS B 224
                                    64.798
                                                     93.589
                                                              1.00
                                                                   20.65
                                             -1.142
ATOM
        4784
               N
                                    64.755
                                                              1.00 27.00
                                             0.311
                                                     93.462
        4785
               CA_
                   LYS B 224
MOTA
                                                              1.00 36:47
                                    64.577
                   LYS B 224
                                              0.938
                                                     94.844
        4786
               CB
MOTA
                                              0.389
                                                      95.-651
                                                              1.00
                                                                    37.72
                   LYS .B 224
                                    63.415
ATOM
        4787
               CG
                                             0.833
                                                     97.101
                                                              1.00 42.06
                                    63.541
               CD
                   LYS B 224
MOTA
        4788
                                                              1.00 45.18
                                              0.276
                                                      97.955
        4789
                   LYS B 224
                                    62.420
               CE
ATOM
                                    62.645
                                              0.570
                                                      99.399
                                                              1.00 46.30
                   LYS B 224
MOTA
        4790
               NZ
                                                              1.00 27.01
                                    66.071
                                              0.808
                                                      92.874
        4791
                   LYS B 224
MOTA
                                                     92.995
                                                              1.00 21.54
                                    67.098
                                              0.139
                   LYS B 224
        4792
               0
ATOM
                                                              1.00 22.66
                   GLY B 225
                                    66.038
                                              1.989
                                                     92.259
MOTA
        4793
               N
                                              2.565
                                                      91.669
                                                              1.00 25.67
                                    67.239
                   GLY B 225
MOTA
        4794
               CA
                                             1.809
                                                     90.459
                                    67.768
                                                               1.00
                                                                    24.95
                   GLY B 225
        4795
               С
MOTA
                                                              1.00
                                                                    26.83
                   GLY B 225
                                    68.917
                                              1.975
                                                      90.069
               ö
ATOM
        4796
                                              0.980
                                                      89.855
                                                              1.00 21.79
                                    66.926
                   LEU B 226
        4797
              N
MOTA
                                              0.180
                                    67.319
                                                      88.692
                                                               1.00
                                                                    22.31
                   LEU B 226
        4798
               CA
ATOM
                                                              1,00 23.29
                                                      88.099
                                             -0.473
                   LEU B 226
                                    66.067
               CB
 MOTA
        4799
                                                               1.00 26.71
                                    66.238
                                             -1.605
                                                      87.091
                   LEU B 226
ATOM
        4800
               CG
                                                      87.813
                                                               1.00
                                                                    26.44
                   LEU B 226
                                    66.846
                                             -2.804
MOTA
        4801
               CD1
                                    64.877
                                             -1.997
                                                      86.508
                                                               1.00 22.96
                   LEU B
                          226
        4802
               CD2
 ATOM
                                                               1.00 22.66
                                    68.008
                                              1.017
                                                      87.603
                   LEU B 226
        4803
               C
 MOTA
                                                                    20.19
                                    67.517
                                              2.087
                                                      87.250
                                                               1.00
                   LEU B 226
 MOTA
        4804
               0
                                    69.134
                                              0.549
                                                      87.060
                                                               1.00 15.52
        4805
               N
                   ASN B 227
 MOTA
                                                      85.998
                                                               1.00
                                                                    19.49
                                    69.794
                                              1.317
                   ASN B 227
MOTA
        4806
               CA
                                                               1.00 20.43
                   ASN B 227
                                    71.304
                                              1.474
                                                      86,270
        4807
               CB
 ATOM
                                                               1.00 28.97
                   ASN B 227
                                    72.062
                                              0.161
                                                      86.206
 ATOM
        4808
               CG
                                    72.015
                                             -0.546
                                                      85.199
                                                               1.00
                                                                    24.30
        4809
               OD1
                   ASN B 227
MOTA
                                                      87.276
                                                               1.00
                                                                    20.88
                                    72.786
                                             -0.160
        4810
              ND2
                   ASN B 227
 ATOM
                                                               1.00 21.26
                   ASN B 227
                                    69.548
                                              0.671
                                                      84.630
 MOTA
        4811
               C
                                                                    18.90
                                             -0.432
                                                      84.555
                                                               1.00
                                    69.004
                   ASN B 227
 ATOM
        4812
               0
                                    69.949
69.720
                                                               1.00 20.98
                                              1.347
                                                      83.552
 MOTA
        4813
              N
                   ASP B
                          228
                                                               1.00
                                                                    22.61
                                                      82.208
              CA
                   ASP B 228
                                              0.817
        4814
 MOTA
                                                               1.00 23.46
                                    70.270
                                              1.753
                                                      31.126
                   ASP B 228
               CB
 ATOM
        4815
                                    69.596
                                              3.113
                                                      81.119
                                                               1.00 26.12
 MOTA
        4816
               CG
                   ASP 3 228
                                                               1.00
                                                                    26.75
                                              3.193
                                                      81.415
        4817
               OD1
                   ASP B -228
                                    68.387
 ATOM
                                                      80.773
                                                               1.00 30.22
                   ASP B 228
                                    70.276
                                              4.101
               OD2
        4818
 ATOM
                                                               1.00 23.49
                                                      81.952
                                    70.286
                                             -0.573
                   ASP B 228
 MOTA
        4819
               C
                                                      81.288
                                                               1.00 19.31
                                             -1.390
                                    69.651
        4820
                   ASP B 228
               0
 MOTA
                                                      82.453
                                                               1.00 22.24
                   ASN B 229
                                    71.484
                                             -0.836
              N
        4821
 ATCM
                                                               1.00 23.30
                                                      82.250
                                    72.111
                                             -2.135
                   ASN B 229
 ATOM
        4822
              CA
                                                      82.737
                                                               1.00 20.99
                                    73.562
                                             -2.101
        4823
              CB
                   ASN B 229
 ATOM
                                             -1.237
                                                               1.00 25.71
                                                      81.859
                                    74.441
        4824
               CG
                   ASN B 229
 ATOM
                                                               1.00 26.40
                                                      80.678
                                    74.644
                                             -1.538
                   ASN B 229
 atom
        4825
              OD1
                                                               1.00 27.44
                                                      82.417
                                    74.955
                                             -0.151
        4826
              ND2
                   ASN B 229
 ATOM
                                             -3.252
                                                      82.943
                                                               1.00 23.74
                                    71.341
                   ASN B 229
        4827
               С
 ATOM
                                                      82.402
                                                               1.00 20.51
                                    71.207
                                             -4.346
                   ASN B 229
               0
 ATOM
        4828
                                                               1.00 23.06
                                    70.832
                                             -2.976
                                                      84.139
        4829
              N
                   GLU B 230
 ATOM
                                                               1.00 23.01
                                             -3.977
                                                      84.874
                                    70.069
              CA
                   GLU B
                          230
        4830
 MOTA
                                                               1.00 24.73
                                                      86.297
                                    69.799
                                             -3.480
              СВ
                   GLU B 230
        4831
 ATOM
                                                               1.00 27.28
                                                      87.087
                                    71.069
                                             -3.201
                   GLU B 230
 MOTA
        4832
              CG
                                                               1.00 27.47
                                    70.792
                                                      88.470
                                             -2.649
               CD
                   GLU B
                         230
        4833
 MOTA
                                                      88.569
                                                               1.00 27.87
                                    70.086
                                             -1.625
                  GLU B 230
              OE1
 MOTA
        4834
                                                                    26.79
                                                      89.455
                                    71.286
                                             -3.232
                                                               1.00
                   GLU B 230
        4835
              OE2
 ATOM
                                                               1.00 24.25
                                                      84.146
                                    68.749
                                             -4.281
                   GLU B
                         230
        4836
               С
 ATOM
                                             -5.445
                                                      84.022
                                                               1.00 15.89
                   GLU B 230
                                    68.347
 ATCM
        4837
              0
                                                               1.00 21.46
                                    68.091
                                             -3.242
                                                      83.637
                   PHE B 231
        :838
              N
 ATCM
                                                               1.00 22.84
                                                      82.933
                                    66.814
                                             -3.429
        4839
              CA
                   PHE B
                         231
 ATOM
                                             -2.079
                                                               1.00 23.96
                                    66.210
                                                      82.529
                   PHE B 231
              CB
        4840
 ATOM
                                                               1.00 26.13
                                                      81.975
                                    64.803
                                             -2.182
        4841
                   PHE B 231
              CG
 ATOM
```

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AROM 4843 CD2 PHE B 231 64.550 -1.956 80.627 1.00 22.9 AROM 4844 CEI PHE B 231 63.250 -2.059 80.114 1.00 25.0 AROM 4845 CEZ PHE B 231 63.250 -2.059 80.114 1.00 25.0 AROM 4846 CZ PHE B 231 66.271 -5.239 80.057 1.00 22.9 AROM 4847 C PHE B 231 66.271 -5.239 80.057 1.00 22.9 AROM 4848 O PHE B 231 66.271 -5.239 80.057 1.00 22.0 AROM 4849 N LEU B 232 67.963 -3.952 80.845 1.00 22.0 AROM 4850 CA LEU B 232 68.200 -4.697 79.614 1.00 19.5 AROM 4851 CB LEU B 232 68.200 -4.697 79.614 1.00 19.5 AROM 4851 CB LEU B 232 68.665 -2.581 78.263 1.00 24.5 AROM 4852 CG LEU B 232 68.665 -2.581 78.263 1.00 24.5 AROM 4855 CD LEU B 232 68.665 -2.581 78.263 1.00 24.5 AROM 4855 CD LEU B 232 68.665 -7.051 79.162 1.00 19.2 AROM 4855 CD LEU B 232 68.686 6-1197 9.998 1.00 19.2 AROM 4855 CD LEU B 232 68.686 6-1197 9.998 1.00 19.2 AROM 4855 CD LEU B 232 68.686 6-1.197 9.998 1.00 19.2 AROM 4856 CD LEU B 233 69.500 -7.599 81.378 1.00 20.7 AROM 4856 CD PHE B 233 70.825 -7.471 1.2 2.632 1.00 19.2 AROM 4856 CD PHE B 233 70.825 -7.471 1.2 2.632 1.00 19.2 AROM 4856 CD PHE B 233 70.825 -7.471 82.632 1.00 20.5 AROM 4850 CD PHE B 233 70.825 -7.471 82.632 1.00 20.5 AROM 4850 CD PHE B 233 70.825 -7.471 82.632 1.00 20.5 AROM 4850 CD PHE B 233 70.825 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.825 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.825 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -9.519 81.378 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -7.471 82.632 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -7.471 83.262 1.00 20.5 AROM 4860 CD PHE B 233 70.827 -7.471 83.262 1.00 20.5 AROM 4861 CD PHE B 233 70.807 -7.599 81.378 1.00 20.6 AROM 4862 CD PHE B 233 70.807 -7.599 81.378 1.00 20.6 AROM 4863 CEI PHE B 233 70.807 -7.599 81.378 1.00 20.6 AROM 4860 CD PHE B 233 70.807 -7.599 81.379 1.00 20.6 AROM 4860 CD PHE B 233 70.807 -7.599 81.00 2.70 1.00 20.6 AR									1 00 05 00
AROM 4843 CD2 PHE B 2311 62.450 -1.956 80.627 1.00 22.9 AROM 4845 CE2 PHE B 2311 62.440 -2.618 82.304 1.00 25.0 AROM 4846 CZ PHE B 2311 62.406 -2.618 82.304 1.00 25.0 AROM 4846 CZ PHE B 2311 62.196 -2.390 80.114 1.00 27.4 AROM 4846 CZ PHE B 2311 66.297 -2.398 80.197 1.00 20.2 AROM 4849 N LEU B 231 66.978 -4.288 81.677 1.00 20.1 AROM 4849 N LEU B 232 67.963 -3.952 80.845 1.00 20.0 AROM 4851 CB LEU B 232 68.200 -4.697 79.614 1.00 29.0 AROM 4851 CB LEU B 232 68.665 -2.581 78.263 1.00 29.7 AROM 4851 CB LEU B 232 69.746 -1.856 77.454 1.00 24.5 AROM 4853 CD1 LEU B 232 69.746 -1.856 77.454 1.00 28.1 AROM 4855 C LEU B 232 68.665 -7.951 78.263 1.00 29.7 AROM 4855 C LEU B 232 68.665 -7.051 79.162 1.00 19.9 AROM 4857 N PHE B 233 69.468 -6.119 79.898 1.00 19.2 AROM 4859 CA PHE B 233 69.950 -7.599 81.378 1.00 20.2 AROM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4850 CG PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4851 CB LEU B 232 68.365 -7.051 79.162 1.00 19.6 AROM 4850 CB PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4850 CB PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4851 CB PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4861 CD PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4861 CD PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4862 CD PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4863 CR PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4864 CE2 PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4865 CR PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4866 CR PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4867 CD PHE B 233 70.825 -7.471 82.632 1.00 20.2 AROM 4868 N ALA B 234 67.822 -9.519 83.239 1.00 28.2 AROM 4868 N ALA B 234 67.822 -9.519 83.250 1.00 20.2 AROM 4867 CD PHE B 233 70.825 -9.519 83.250 1.00 20.2 AROM 4868 N ALA B 234 66.826 -8.808 81.779 1.00 20.2 AROM 4868 N ALA B 234 67.822 -9.519 82.550 1.00 20.2 AROM 4869 C R PHE B 233 70.825 -9.519 83.174 1.00 20.2 AROM 4869 N E R PHE B 233 70.825 -9.519 83.174 1.00 20.2 AROM 4870 C B LEU B 235 63.66 -7.288 83.174 1.00 20.2 AROM 48	MΩTrΔ	4842	CD1	.PHE I	3 231	63.738	-2.514		1.00 25.00
ARTOM 4844 CEL PHE B 231						64.550	-1.956	80.627	1.00 22.93
ATOM 4845 CE2 PHE B 231 63.750 -2.059 80.114 1.00 27.4 ATOM 4846 CZ PHE B 231 66.978 -4.288 81.677 1.00 20.2 ATOM 4848 O PHE B 231 66.978 -4.288 81.677 1.00 20.2 ATOM 4848 O PHE B 231 66.978 -4.288 81.677 1.00 22.0 ATOM 4850 CA LEU B 232 67.963 -3.952 80.845 1.00 22.0 ATOM 4851 CB LEU B 232 67.963 -3.952 80.845 1.00 22.0 ATOM 4851 CB LEU B 232 69.192 -3.942 78.734 1.00 24.5 ATOM 4853 CDL LEU B 232 69.192 -3.942 78.734 1.00 24.5 ATOM 4853 CDL LEU B 232 69.192 -3.942 77.814 1.00 29.7 ATOM 4854 CD2 LEU B 232 69.468 -6.180 77.454 1.00 28.1 ATOM 4855 CD LEU B 232 68.665 -2.581 78.263 1.00 29.7 ATOM 4855 CD LEU B 232 68.365 -7.051 79.162 1.00 19.6 ATOM 4856 CD LEU B 232 68.365 -7.051 79.162 1.00 19.6 ATOM 4856 CD LEU B 232 69.468 -6.180 80.962 1.00 20.2 ATOM 4855 CD PHE B 233 69.468 -6.180 80.962 1.00 20.2 ATOM 4856 CD LEU B 233 69.468 -6.180 80.962 1.00 20.2 ATOM 4856 CD LEU B 233 70.825 -7.559 81.378 1.00 20.7 ATOM 4866 CD PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4866 CD PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4866 CD PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4866 CD PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4866 CD PHE B 233 70.815 -9.328 84.294 1.00 25.3 ATOM 4866 CD PHE B 233 70.815 -9.328 84.294 1.00 25.3 ATOM 4866 CD PHE B 233 70.811 -9.328 84.294 1.00 25.3 ATOM 4866 CD PHE B 233 70.81 -9.328 84.294 1.00 25.3 ATOM 4867 CD PHE B 233 70.81 -9.328 84.294 1.00 25.3 ATOM 4868 N ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4868 N ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4868 N ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4868 N ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4868 N ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4870 CD ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4870 CD ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4871 CD ALA B 234 65.853 -9.567 81.270 1.00 20.3 ATOM 4878 CD LEU B 235 66.865 -9.9197 76.644 1.00 21.4 ATOM 4878 CD LEU B 235 66.865 -9.9197 76.644 1.00 21.4 ATOM 4879 CD ALA B 234 65.275 -9.00 77.599 1.00 24.4 ATOM 4889 N CALUB S 235 66.865 -9.9197 76.548 1.00 2							-2.618	82.304	1.00 25.03
AROM								80.114	1.00 27.46
AROM 4848 O PHE B 231 66.978 -4.288 81.677 1.00 23.1 AROM 4848 O PHE B 231 66.221 -5.239 81.464 1.00 20.2 C AROM 4850 CA LEU B 232 67.963 -3.952 80.845 1.00 22.0 C AROM 4851 CB LEU B 232 67.963 -3.952 80.845 1.00 22.0 C AROM 4851 CB LEU B 232 69.192 -3.942 78.734 1.00 24.5 AROM 4853 CD1 LEU B 232 69.192 -3.942 78.734 1.00 24.5 AROM 4853 CD1 LEU B 232 69.468 65 -2.581 78.263 1.00 29.7 AROM 4854 CD2 LEU B 232 69.468 -6.187 79.898 10.01 19.5 AROM 4855 C LEU B 232 68.665 -7.051 79.162 1.00 20.7 AROM 4856 C LEU B 232 68.365 -7.051 79.162 1.00 19.6 AROM 4857 N PHE B 233 69.468 -6.180 80.962 1.00 20.7 AROM 4856 C PHE B 233 69.468 -6.280 80.962 1.00 20.7 AROM 4856 C PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4861 CD1 PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4863 CE1 PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4863 CE1 PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4863 CE1 PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4863 CE1 PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4866 C PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4866 C PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4866 C PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4866 C PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4867 O PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4867 O PHE B 233 70.825 -7.471 82.632 1.00 23.7 AROM 4867 O PHE B 233 70.83 -10.573 84.832 1.00 21.7 AROM 4867 O PHE B 233 70.83 -10.573 84.832 1.00 21.7 AROM 4867 O PHE B 233 70.83 -10.573 84.832 1.00 21.7 AROM 4867 C PHE B 233 70.83 -10.573 84.832 1.00 21.7 AROM 4867 O PHE B 233 66.553 -9.567 81.270 1.00 22.7 AROM 4869 C A ALA B 234 66.525 -8.576 82.963 1.00 19.4 AROM 4867 O PHE B 233 66.553 -9.567 81.270 1.00 22.7 AROM 4870 C B ALA B 234 66.525 -8.576 82.963 1.00 19.4 AROM 4871 C ALA B 234 66.525 -8.576 82.963 1.00 19.4 AROM 4875 C B ALA B 234 66.525 -7.793 80.950 1.00 19.4 AROM 4875 C B ALA B 234 66.525 -7.793 80.950 1.00 19.4 AROM 4875 C B ALA B 234 66.525 -7.793 80.950 1.00 19.4 AROM 4879 C ALA B 234 66.525 -7.793 80.950 1.00 19.4 AROM 4870 C B ALA B 234 66.525 -7.793 80.950 1	MOTA								
AROM A848									
MYOM 4848 N 100 BEB B 231 ATOM 4850 CA 100 BEU B 232 67.963 - 3.952 ATOM 4851 CB 4851 CB 100 BEU B 232 68.200 - 4.697 78.734 1.00 24.5 ATOM 4851 CB 101 EU B 232 68.200 - 4.697 78.734 1.00 24.5 ATOM 4852 CG 102 BEU B 232 68.665 - 2.581 78.263 1.00 29.7 ATOM 4854 CD2 102 LEU B 232 68.665 - 7.051 79.162 1.00 19.5 ATOM 4855 CD 102 LEU B 232 68.665 - 7.051 79.162 1.00 19.5 ATOM 4856 CD 102 BEU B 232 68.665 - 7.051 79.162 1.00 19.5 ATOM 4857 N PHE B 233 69.468 6.280 69.962 1.00 20.5 ATOM 4858 CA ATOM 4858 CA 100 PHE B 233 69.950 - 7.599 81.378 1.00 20.5 ATOM 4861 CD1 4861 CD1 4862 CD2 4863 CC2 4864 CC2 4864 CC2 4864 CC2 4864 CC2 4864 CC2 4865 CA 4866 CD2 4866 CC3 4866 CC3 4866 CC3 4866 CC4 4867 CO4 4867 CO4 4868 CA 4868 CA 4868 CA 4868 CA 4868 CA 4868 CA 4869 CA 4869 CA 4860 CC9 4868 CA 4860 CC9	MOTA	4847	С						
ARTOM 4849 N LEU B 232 67.963 -3.952 80.845 1.00 22.0 ARTOM 4850 CA LEU B 232 69.192 -3.942 78.734 1.00 24.5 ARTOM 4851 CB LEU B 232 69.192 -3.942 78.734 1.00 24.5 ARTOM 4853 CD LEU B 232 69.746 -1.856 77.454 1.00 24.5 ARTOM 4853 CD LEU B 232 69.746 -1.856 77.454 1.00 26.5 ARTOM 4855 C LEU B 232 69.746 -1.856 77.454 1.00 26.5 ARTOM 4856 C LEU B 232 68.365 -7.051 79.162 1.00 19.5 ARTOM 4857 N PHE B 233 69.468 -6.220 80.962 1.00 20.7 ARTOM 4858 CA PHE B 233 69.468 -6.280 80.962 1.00 20.7 ARTOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 23.7 ARTOM 4850 CG PHE B 233 70.825 -7.471 82.632 1.00 23.7 ARTOM 4861 CD1 PHE B 233 70.825 -9.519 82.731 1.00 30.7 ARTOM 4863 CE1 PHE B 233 70.825 -9.519 82.731 1.00 30.7 ARTOM 4863 CE1 PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4866 CC PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4866 CC PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4866 CC PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4866 CC PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4866 CC PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4866 CC PHE B 233 70.881 -9.328 84.294 1.00 25.3 ARTOM 4867 OPHE B 233 68.712 -8.439 81.727 1.00 20.7 ARTOM 4867 OPHE B 233 68.712 -8.439 81.727 1.00 20.7 ARTOM 4867 OPHE B 233 68.553 -9.567 81.270 1.00 21.3 ARTOM 4867 OPHE B 233 66.535 -7.733 82.963 1.00 19.4 ARTOM 4870 CB ALA B 234 66.265 -7.733 82.963 1.00 19.4 ARTOM 4871 C ALA B 234 66.626 -7.878 82.560 1.00 21.3 ARTOM 4870 CB ALA B 234 66.626 -7.878 82.560 1.00 21.3 ARTOM 4871 C ALA B 234 66.626 -7.878 82.560 1.00 21.3 ARTOM 4871 C ALA B 234 66.626 -9.576 82.963 1.00 19.4 ARTOM 4876 CB LEU B 235 64.872 -7.878 82.560 1.00 21.3 ARTOM 4876 CB LEU B 235 64.872 -7.878 82.560 1.00 21.3 ARTOM 4876 CB LEU B 235 64.872 -7.877 77.664 1.00 20.7 ARTOM 4878 CD LEU B 235 64.872 -7.878 82.560 1.00 21.3 ARTOM 4878 CD LEU B 235 63.465 -7.733 83.950 1.00 19.4 ARTOM 4878 CD LEU B 235 64.872 -7.877 77.664 1.00 20.7 ARTOM 4880 C G LEU B 235 66.666 -9.191 7.8431 1.00 30.2 ARTOM 4880 C G LEU B 235 66.666 -9.191 7.8431 1.00 30.2 ARTOM 4881 CD GLU B 2		4848	0	PHE I	B 231	66.221	-5.239		
ATOM 4850 CA LEU B 232 68.200 -4.697 79.614 1.00 19.5 ATOM 4851 CB LEU B 232 69.746 -1.856 77.454 1.00 24.5 ATOM 4853 CD1 LEU B 232 69.746 -1.856 77.454 1.00 24.5 ATOM 4854 CD2 LEU B 232 69.746 -1.856 77.454 1.00 26.5 ATOM 4855 C LEU B 232 68.665 -2.581 78.263 1.00 29.7 ATOM 4855 C LEU B 232 68.668 -6.119 79.898 1.00 19.2 ATOM 4856 O LEU B 232 68.668 -6.119 79.898 1.00 19.2 ATOM 4857 N PHE B 233 69.468 -6.280 80.962 1.00 20.5 ATOM 4857 N PHE B 233 69.468 -6.280 80.962 1.00 20.5 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 20.5 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 20.5 ATOM 4850 CC PHE B 233 70.825 -9.519 82.731 1.00 30.4 ATOM 4861 CD1 PHE B 233 72.285 -9.519 82.731 1.00 30.4 ATOM 4863 CE2 PHE B 233 70.803 -10.573 84.294 1.00 25.3 ATOM 4864 CE2 PHE B 233 70.803 -10.573 84.294 1.00 25.3 ATOM 4866 C PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4867 C PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4868 C PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4868 C PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4867 C PHE B 233 68.553 -9.567 81.270 1.00 20.5 ATOM 4867 C PHE B 233 68.553 -9.567 81.270 1.00 20.5 ATOM 4867 C PHE B 233 68.553 -9.567 81.270 1.00 21.4 ATOM 4869 CA ALA B 234 65.855 -7.733 83.950 1.00 19.3 ATOM 4867 C PHE B 233 68.553 -9.567 81.270 1.00 21.4 ATOM 4870 C B ALA B 234 65.835 -7.733 83.950 1.00 19.3 ATOM 4867 C PHE B 235 66.656 8.595 -7.733 83.950 1.00 19.3 ATOM 4870 C B ALA B 234 65.835 -7.733 83.950 1.00 19.3 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.3 ATOM 4871 C ALA B 234 65.725 -7.79 87.95 1.00 19.3 ATOM 4871 C ALA B 234 65.725 -7.79 87.95 1.00 19.3 ATOM 4873 N LEU B 235 66.656 -7.79 77.3 80.815 1.00 20.7 ATOM 4873 N LEU B 235 66.665 -9.91 78.79 77.664 1.00 27.4 ATOM 4873 N LEU B 235 66.665 -9.91 78.79 77.664 1.00 27.4 ATOM 4870 C B ALA B 234 65.770 -9.276 78.795 1.00 24.4 ATOM 4870 C B ALA B 234 65.770 -9.276 78.795 1.00 24.4 ATOM 4880 O LEU B 235 66.666 6.665 -9.91 78.795 77.094 1.00 27.4 ATOM 4880 O LEU B 235 66.660 6.657 79.09 77.659 1.00 29.4 ATOM 4880 O LEU B 235						67.963	-3.952	80.845	1.00 22.02
ATOM 4851 CB LEU B 232 69.192 -3.942 78.734 1.00 24.5 ATOM 4853 CD1 LEU B 232 68.665 -2.581 78.263 1.00 29.5 ATOM 4854 CD2 LEU B 232 67.409 -2.784 77.414 1.00 26.5 ATOM 4855 C LEU B 232 68.665 -7.051 79.162 1.00 19.2 ATOM 4856 O LEU B 232 68.365 -7.051 79.162 1.00 19.2 ATOM 4857 N PHE B 233 69.365 -7.051 79.162 1.00 19.2 ATOM 4858 CA PHE B 233 69.950 -7.599 81.378 1.00 20.5 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 20.5 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 23.4 ATOM 4850 CB PHE B 233 70.825 -7.471 82.632 1.00 23.4 ATOM 4861 CD1 PHE B 233 70.825 -7.599 81.378 1.00 20.5 ATOM 4863 CEI PHE B 233 70.825 -7.599 81.378 1.00 20.5 ATOM 4863 CEI PHE B 233 70.825 -7.571 82.632 1.00 25.4 ATOM 4863 CEI PHE B 233 70.825 -7.573 84.832 1.00 30.4 ATOM 4863 CEI PHE B 233 70.825 -7.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 70.803 -10.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 70.881 -9.528 84.234 1.00 25.3 ATOM 4866 C PHE B 233 70.835 -7.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 70.835 -7.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 70.835 -7.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 70.835 -7.573 84.832 1.00 31.6 ATOM 4867 O PHE B 233 68.553 -9.567 81.270 1.00 21.3 ATOM 4868 N ALA B 234 66.626 -8.576 82.963 1.00 21.4 ATOM 4870 CB ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 65.835 -7.733 89.950 1.00 19.4 ATOM 4871 C B ALA B 234 65.835 -7.733 89.950 1.00 19.4 ATOM 4874 CA LEU B 235 66.485 -7.733 89.950 1.00 19.4 ATOM 4876 CG LEU B 235 64.877 -8.898 81.749 1.00 18.4 ATOM 4876 CG LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4876 CG LEU B 235 66.666 -9.9191 78.462 1.00 21.4 ATOM 4878 CD LEU B 235 66.666 -9.9191 78.462 1.00 21.4 ATOM 4883 CG LEU B 235 66.666 -9.9191 78.462 1.00 21.4 ATOM 4880 C LEU B 235 66.666 -9.9191 78.462 1.00 21.4 ATOM 4881 N GLU B 236 66.666 -9.9191 78.462 1.00 22.4 ATOM 4880 C LEU B 235 66.666 -9.9191 78.462 1.00 22.4 ATOM 4880 C C LEU B 235 66.666 -9.9191 78.462 1.00 22.4 ATOM 4880 C C LEU B 235 66.666 -9.9191 78.462 1.00 22.4 ATOM 4880 C C LEU B 235 66.666 -9.9191 78.4							-4.697	79.614	1.00 19.97
ATOM 4852 CG LEU B 232 68.665 -2.581 78.263 1.00 29.7 ATOM 4853 CD1 LEU B 232 69.746 -1.856 77.444 1.00 26.5 ATOM 4854 CD2 LEU B 232 67.409 -2.784 77.414 1.00 26.5 ATOM 4855 C LEU B 232 68.688 -6.119 79.898 1.00 19.2 ATOM 4856 N PHE B 233 69.468 -6.280 80.962 1.00 20.5 ATOM 4857 N PHE B 233 69.468 -6.280 80.962 1.00 20.5 ATOM 4858 CA PHE B 233 70.825 -7.959 81.378 1.00 20.5 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 23.7 ATOM 4860 CG PHE B 233 70.825 -7.471 82.632 1.00 23.7 ATOM 4861 CD1 PHE B 233 70.825 -7.959 81.378 1.00 20.3 ATOM 4862 CD2 PHE B 233 70.481 -9.328 84.294 1.00 25.7 ATOM 4863 CE1 PHE B 233 70.481 -9.328 84.294 1.00 25.7 ATOM 4864 CE2 PHE B 233 70.481 -9.328 84.294 1.00 25.7 ATOM 4865 C PHE B 233 70.811 -10.22 84.317 1.00 20.7 ATOM 4866 C PHE B 233 70.811 -11.292 84.317 1.00 20.7 ATOM 4867 O PHE B 233 68.712 -8.439 81.770 1.00 20.7 ATOM 4867 O PHE B 233 68.712 -8.439 81.770 1.00 20.7 ATOM 4867 O PHE B 233 68.712 -8.439 81.770 1.00 20.7 ATOM 4867 O PHE B 233 68.513 -9.567 81.270 1.00 21.7 ATOM 4867 O PHE B 233 68.513 -9.567 81.270 1.00 21.7 ATOM 4867 O PHE B 233 68.513 -9.567 81.270 1.00 21.7 ATOM 4867 O PHE B 233 68.513 -9.567 81.270 1.00 21.7 ATOM 4869 CA ALA B 234 65.825 -7.878 82.560 1.00 19.4 ATOM 4870 CB ALA B 234 65.825 -7.878 82.560 1.00 19.4 ATOM 4871 C ALA B 234 65.8772 -8.898 81.749 1.00 18.4 ATOM 4874 CA LEU B 235 65.634 -7.934 80.845 1.00 20.7 ATOM 4876 CG LEU B 235 66.772 -8.898 81.749 1.00 18.4 ATOM 4876 CG LEU B 235 66.867 -7.997 77.664 1.00 26.7 ATOM 4876 CG LEU B 235 66.771 -9.897 77.664 1.00 26.7 ATOM 4870 CB ALA B 234 65.825 -7.70 1.00 20.7 ATOM 4870 CB ALA B 234 65.855 -7.70 1.00 20.7 ATOM 4870 CB ALA B 234 65.855 -7.70 1.00 20.7 ATOM 4870 CB ALA B 234 65.855 -7.70 1.00 20.7 ATOM 4870 CB ALA B 234 65.855 -7.70 1.00 20.7 ATOM 4870 CB ALA B 235 66.665 -9.791 77.664 1.00 26.7 ATOM 4870 CD ALA B 235 66.665 -9.791 77.664 1.00 26.7 ATOM 4870 CB ALA B 235 66.70 1.00 81.645 1.00 20.7 ATOM 4870 CB ALA B 236 67.00 -9.00 77.629 1.00 20.7 ATOM 4880 CD ALEU B 235 66.765 -9.7997 77.66									1.00 24.99
ATOM 4853 CD1 LEU B 232 69.746 -1.856 77.454 1.00 28.1 ATOM 4855 C LEU B 232 68.688 -6.119 79.898 1.00 19.2 ATOM 4856 O LEU B 232 68.688 -6.119 79.898 1.00 19.2 ATOM 4857 N PHE B 233 69.950 -7.599 81.378 1.00 20.3 ATOM 4858 CA PHE B 233 70.825 -7.051 79.162 1.00 19.4 ATOM 4858 CA PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4858 CA PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4850 CG PHE B 233 70.825 -7.471 82.632 1.00 20.3 ATOM 4860 CG PHE B 233 70.825 -7.519 81.378 1.00 20.3 ATOM 4861 CD1 PHE B 233 72.885 -9.519 82.731 1.00 30.4 ATOM 4863 CEI PHE B 233 70.885 -9.519 82.731 1.00 30.5 ATOM 4863 CEI PHE B 233 70.881 -9.519 82.731 1.00 32.3 ATOM 4863 CEI PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4864 CE2 PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4867 CP PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4867 CP PHE B 233 68.751 -8.7878 82.560 1.00 21.3 ATOM 4868 N ALAB 8234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALAB 8234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALAB 8234 65.835 -9.567 81.270 1.00 21.3 ATOM 4871 C B ALAB 8234 65.835 -7.733 83.950 1.00 19.4 ATOM 4873 N LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4873 N LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4876 CB LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4877 CD LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 66.482 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 66.868 -9.191 78.462 1.00 20.3 ATOM 4881 N GLU B 235 66.865 -9.191 78.462 1.00 20.3 ATOM 4881 N GLU B 235 66.666 -9.591 77.664 1.00 20.3 ATOM 4880 C GLU B 235 66.666 -9.9191 78.462 1.00 19.4 ATOM 4881 N GLU B 235 66.666 -9.9191 78.462 1.00 19.4 ATOM 4880 C GLU B 235 66.666 -9.9191 78.462 1.00 19.4 ATOM 4880 C GLU B 235 66.666 -9.9191 78.462 1.00 19.4 ATOM 4881 N GLU B 236 67.303 -10.206 77.6991 1.00 24.4 ATOM 4880 C GLU B 236 67.00 -9.032 76.5591 1.00 45.4 ATOM 4880 C GLU B 236 67.00 -9.032 76.5591 1.00 45.4 ATOM 4880 C GLU B 236 67.303 -10.206 77.6991 1	MOTA					•			
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ATOM 4855 CD LEU B 232 67.409 -2.784 77.414 .00 26.24 ATOM 4855 C LEU B 232 68.868 -6.119 79.898 1.00 19.24 ATOM 4856 C LEU B 232 68.868 -6.119 79.898 1.00 19.24 ATOM 4857 N PHE B 233 69.468 -6.280 80.952 1.00 20.25 ATOM 4858 CA PHE B 233 70.825 -7.471 82.632 1.00 20.34 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 20.34 ATOM 4861 CD1 PHE B 233 71.217 -8.790 83.239 1.00 28.54 ATOM 4861 CD1 PHE B 233 72.285 -9.519 82.731 1.00 30.44 ATOM 4863 CE1 PHE B 233 72.285 -9.519 82.731 1.00 30.44 ATOM 4863 CE1 PHE B 233 70.803 -10.573 84.832 1.00 31.44 ATOM 4864 CE2 PHE B 233 70.803 -10.573 84.832 1.00 31.44 ATOM 4866 C PHE B 233 70.803 -10.573 84.832 1.00 31.44 ATOM 4866 C PHE B 233 68.553 -9.567 81.270 1.00 20.34 ATOM 4866 C PHE B 233 68.553 -9.567 81.270 1.00 20.34 ATOM 4868 N ALA B 234 66.626 -8.576 82.960 1.00 21.34 ATOM 4869 CA ALA B 234 66.626 -8.576 82.961 1.00 21.34 ATOM 4867 O PHE B 233 68.712 -8.898 81.772 1.00 20.34 ATOM 4869 CA ALA B 234 65.835 -7.733 83.950 1.00 19.34 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.34 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 21.34 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 21.34 ATOM 4875 CB LEU B 235 66.473 -6.874 78.795 1.00 21.34 ATOM 4876 CG LEU B 235 66.473 -6.874 78.795 1.00 21.34 ATOM 4876 CG LEU B 235 66.4773 -6.874 78.795 1.00 21.34 ATOM 4878 CD LEU B 235 64.648 -10.205 78.431 1.00 19.34 ATOM 4878 CD LEU B 235 63.763 -9.276 78.931 1.00 19.34 ATOM 4882 CA GLU B 235 64.648 -10.205 78.831 1.00 19.34 ATOM 4883 CB GLU B 235 64.648 -10.205 78.831 1.00 19.34 ATOM 4880 N LEU B 235 66.655 -9.8797 77.654 1.00 27.34 ATOM 4880 N LEU B 235 66.665 -9.8797 77.654 1.00 27.34 ATOM 4881 N GLU B 235 66.665 -9.8797 77.654 1.00 27.34 ATOM 4881 N GLU B 235 66.665 -9.8797 77.654 1.00 27.34 ATOM 4880 N LEU B 235 66.6665 -9.8797 77.654 1.00 27.34 ATOM 4880 N LEU B 235 66.6665 -9.8797 77.654 1.00 20.34 ATOM 4880 N LEU B 235 66.6665 -9.8797 77.654 1.00 20.34 ATOM 4880 N LEU B 235 67.7697 -9.8797 77.654 1.00 20.34 ATOM 4880 N LEU B 235 67.7697 -9.932 77.6548 1.00 27.34 ATOM 4		4853	CD1	LEU !	в 232	69.746			
ATOM 4855 C LEU B 232 68.688 -6.119 79.898 1.00 19.4 ATOM 4856 O LEU B 232 68.365 -7.051 79.162 1.00 19.4 ATOM 4857 N PHE B 233 69.950 -7.599 81.378 1.00 20.5 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 23.3 ATOM 4869 CB PHE B 233 70.825 -7.471 82.632 1.00 23.3 ATOM 4860 CG PHE B 233 70.825 -7.471 82.632 1.00 23.3 ATOM 4861 CD1 PHE B 233 70.825 -7.471 82.632 1.00 30.4 ATOM 4862 CD2 PHE B 233 70.81 -9.328 84.294 1.00 25.3 ATOM 4863 CE1 PHE B 233 70.481 -9.328 84.294 1.00 25.3 ATOM 4864 CE2 PHE B 233 70.803 10.573 84.832 1.00 31.4 ATOM 4865 CP PHE B 233 70.803 10.573 84.832 1.00 31.4 ATOM 4866 C PHE B 233 66.555 -9.567 81.270 1.00 20.3 ATOM 4866 C PHE B 233 66.555 -9.567 81.270 1.00 20.3 ATOM 4867 O PHE B 233 66.555 -9.567 81.270 1.00 20.3 ATOM 4869 CA ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.4 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 18.3 ATOM 4876 CB ELEU B 235 64.822 -8.141 79.652 1.00 18.3 ATOM 4877 CD1 LEU B 235 65.634 -7.934 80.845 1.00 20.3 ATOM 4878 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4878 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4880 CA BLEU B 235 64.820 -8.191 78.462 1.00 20.3 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 20.3 ATOM 4880 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 20.3 ATOM 4880 CB LEU B 235 64.648 -10.205 78.231 1.00 1.00 26.3 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.3 ATOM 4880 CB LEU B 235 64.66 655 -9.276 78.024 1.00 27.3 ATOM 4880 CB LEU B 235 64.66 655 -9.276 78.024 1.00 27.3 ATOM 4880 CB LEU B 236 66.665 -9.191 78.462 1.00 24.3 ATOM 4880 CB LEU B 236 67.303 -10.206 77.629 1.00 24.3 ATOM 4880 CB LEU B 236 67.500 -10.207 78.209 1.00 24.3 ATOM 4880 CB LEU B 236 67.70 -9.276 78.311 1.00 26.3 ATOM 4890 N LYS B 237 77.474 1.10.10 18.		4854	CD2	LEU I	B 232	67.409	-2.784	77.414	
ATOM 4857 N PHE B 233 69.468 -6.280 80.962 1.00 19.467 ATOM 4858 CA PHE B 233 69.950 -7.599 81.378 1.00 20.5 ATOM 4858 CA PHE B 233 70.825 -7.471 82.632 1.00 23.7 ATOM 4860 CG PHE B 233 70.825 -7.471 82.632 1.00 23.7 ATOM 4860 CG PHE B 233 70.825 -9.519 82.731 1.00 30.4 ATOM 4861 CD1 PHE B 233 70.825 -9.519 82.731 1.00 30.4 ATOM 4862 CD2 PHE B 233 70.825 -9.519 82.731 1.00 30.4 ATOM 4863 CE1 PHE B 233 70.825 -9.519 82.731 1.00 30.4 ATOM 4864 CE2 PHE B 233 70.803 10.573 84.832 1.00 31.2 ATOM 4865 CD PHE B 233 70.803 10.573 84.832 1.00 31.2 ATOM 4866 C PHE B 233 70.803 10.573 84.832 1.00 31.2 ATOM 4866 C PHE B 233 68.712 -8.439 81.727 1.00 20.3 ATOM 4866 C PHE B 233 68.712 -8.439 81.727 1.00 20.3 ATOM 4867 O PHE B 233 68.712 -8.439 81.727 1.00 21.3 ATOM 4868 N ALA B 234 66.626 -8.576 82.963 1.00 21.3 ATOM 4869 CA ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 65.835 -7.733 83.950 1.00 19.4 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.4 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 21.3 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 21.3 ATOM 4876 CB LEU B 235 63.465 -6.607 78.795 1.00 24.7 ATOM 4876 CB LEU B 235 63.465 -6.607 78.795 1.00 24.7 ATOM 4877 CD LEU B 235 63.465 -6.607 78.795 1.00 24.7 ATOM 4878 CB LEU B 235 63.765 -9.276 78.795 1.00 24.7 ATOM 4878 CB LEU B 235 63.765 -9.276 78.795 1.00 24.7 ATOM 4889 CB LEU B 235 63.766 -9.276 78.795 1.00 24.7 ATOM 4880 CB LEU B 235 63.766 -9.276 78.795 1.00 24.7 ATOM 4881 N GLU B 235 63.766 -9.276 78.795 1.00 24.7 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.7 ATOM 4880 CB LEU B 235 66.866 -11.2.552 77.551 1.00 24.7 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.7 ATOM 4880 CB LEU B 235 66.866 -11.562 77.584 1.00 27.7 ATOM 4880 CB LEU B 235 66.866 -11.607 78.995 1.00 24.7 ATOM 4880 CB LEU B 236 66.665 -9.191 78.462 1.00 24.7 ATOM 4880 CB LEU B 236 66.665 -9.191 78.462 1.00 24.7 ATOM 4880 CB LEU B 236 66.665 -9.191 78.462 1.00 24.7 ATOM 4880 CB LEU B 236 66.665 -9.191 78.462 1.00 24.7 ATOM 4880 CB LEU B 236 66.665 -9.191 78.462 1.00						68.688	-6.119	79.898	1.00 19.25
ATOM 4857 N PHE B 233 69.468 -6.280 80.962 1.00 20.5 ATOM 4858 CA PHE B 233 69.950 -7.599 81.378 1.00 20.5 ATOM 4869 CB PHE B 233 70.825 -7.471 82.632 1.00 23.5 ATOM 4861 CD1 PHE B 233 70.825 -7.471 82.632 1.00 23.5 ATOM 4861 CD1 PHE B 233 70.825 -7.471 82.632 1.00 23.5 ATOM 4862 CD2 PHE B 233 70.825 -7.471 82.632 1.00 30.4 ATOM 4863 CE1 PHE B 233 70.841 -9.328 84.294 1.00 25.5 ATOM 4863 CE1 PHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4864 CE2 PHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4865 CZ PHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4866 CP PHE B 233 66.553 -9.567 81.270 1.00 21.5 ATOM 4867 CO PHE B 233 66.553 -9.567 81.270 1.00 21.5 ATOM 4868 N ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 65.875 -7.733 83.950 1.00 19.4 ATOM 4871 C ALA B 234 65.772 -8.898 81.749 1.00 18.4 ATOM 4871 C ALA B 234 65.253 -10.010 81.624 1.00 21.3 ATOM 4873 N LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4876 CB LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4877 CD1 LEU B 235 63.763 -7.734 80.845 1.00 20.3 ATOM 4878 CD2 LEU B 235 63.763 -9.276 78.024 1.00 27.3 ATOM 4880 CA BLEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4877 CD1 LEU B 235 63.763 -9.276 78.024 1.00 27.3 ATOM 4880 CA BLEU B 235 64.658 -9.191 78.462 1.00 27.3 ATOM 4880 CA BLEU B 235 64.648 -10.205 78.391 1.00 18.3 ATOM 4880 CA BLEU B 235 64.658 -9.191 78.462 1.00 19.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4880 CA BLEU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA BLEU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA BLU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA BLU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA BLU B 236 67.70 -9.276 78.734 1.00 24.4 ATOM 4880 CA BLU B 236 67.70 -9.276 78.734 1.00 24.4 ATOM 4880 CA BLU B 236 67.70 -9.276 78.734 1.00 31.4 ATOM 4880 CA BLU B 236 67.70 -9.276 78.734 1.00 31.4 ATOM 4880 CA BLU B 236 67.70 -9.276 78.731 1.00 24.4 ATOM 4890 N LYS B 237 77.679 98.59 77.501							-7.051	79.162	1.00 19.49
ATOM 4859 CB PHE B 233 70.825 -7.599 81.378 1.00 20.3 ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 23.3 ATOM 4860 CG PHE B 233 70.825 -7.471 82.632 1.00 23.3 ATOM 4861 CD1 PHE B 233 72.285 -9.519 82.731 1.00 30.4 ATOM 4862 CD2 PHE B 233 72.617 -10.762 83.262 1.00 31.4 ATOM 4863 CE1 PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4864 CE2 PHE B 233 70.803 -10.573 84.832 1.00 31.4 ATOM 4865 CP PHE B 233 71.871 -11.292 84.317 1.00 20.3 ATOM 4866 C PHE B 233 71.871 -11.292 84.317 1.00 20.3 ATOM 4867 O PHE B 233 68.712 -8.439 81.727 1.00 20.3 ATOM 4868 N ALA B 234 66.626 -8.576 82.963 1.00 21.4 ATOM 4869 N ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 65.573 -9.567 81.270 1.00 20.3 ATOM 4870 CB ALA B 234 65.573 -9.567 81.749 1.00 18.4 ATOM 4871 C ALA B 234 65.575 -7.733 83.950 1.00 19.4 ATOM 4873 N LEU B 235 66.634 -7.934 80.845 1.00 19.4 ATOM 4874 CA LEU B 235 66.634 -7.934 80.845 1.00 21.4 ATOM 4875 CB LEU B 235 63.465 -6.607 78.024 1.00 21.4 ATOM 4876 CG LEU B 235 63.465 -6.607 78.024 1.00 21.4 ATOM 4877 CD1 LEU B 235 63.465 -6.607 78.024 1.00 20.4 ATOM 4878 CD2 LEU B 235 63.465 -6.607 78.024 1.00 20.4 ATOM 4887 CB LEU B 235 63.465 -6.607 78.024 1.00 20.4 ATOM 4880 C LEU B 235 63.465 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 235 63.763 -9.276 78.795 1.00 20.4 ATOM 4880 C LEU B 235 63.763 -7.833 77.864 1.00 20.4 ATOM 4880 C LEU B 235 66.665 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 235 66.666 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 235 66.666 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 236 67.00 -9.276 78.795 1.00 20.4 ATOM 4880 C LEU B 236 67.80 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 236 67.90 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 236 67.90 -9.191 78.462 1.00 19.4 ATOM 4881 C G GLU B 236 67.90 -9.191 78.462 1.00 19.4 ATOM 4880 C LEU B 236 67.90 -9.191 78.462 1.00 19.4 ATOM 4881 C G GLU B 236 67.00 -9.191 78.462 1.00 19.4 ATOM 4880 C SER B 237 67.449 -13.058 80.130 1.00 27.4 ATOM 4890 N LYS B 237 67.449 -13.058 80.130 1.00 27.4 ATOM 4891 C C LYS B 237 67.449 -13.058 80.931 1.00 39.4 ATO									1.00 20.50
ATOM 4859 CB PHE B 233 70.825 -7.471 82.632 1.00 23.5 ATOM 4860 CG PHE B 233 71.217 -8.790 83.239 1.00 28.5 ATOM 4861 CD1 PHE B 233 70.481 -9.519 82.731 1.00 30.4 ATOM 4862 CD2 PHE B 233 70.481 -9.328 84.294 1.00 25.5 ATOM 4863 CE1 PHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4864 CE2 PHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4865 CZ PHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4866 CPHE B 233 70.803 -10.573 84.832 1.00 31.5 ATOM 4867 OPHE B 233 66.512 -8.439 81.727 1.00 20.5 ATOM 4868 N ALA B 234 66.626 -8.576 82.560 1.00 21.5 ATOM 4869 CA ALA B 234 66.626 -8.576 82.963 1.00 19.6 ATOM 4870 CB ALA B 234 65.835 -7.733 83.950 1.00 19.6 ATOM 4871 C ALA B 234 65.253 -10.010 81.624 1.00 21.5 ATOM 4873 N LEU B 235 66.822 -8.141 79.652 1.00 18.3 ATOM 4875 CB LEU B 235 66.822 -8.141 79.652 1.00 19.4 ATOM 4876 CG LEU B 235 66.822 -8.141 79.652 1.00 19.4 ATOM 4877 CD1 LEU B 235 63.465 -6.607 78.795 1.00 27.4 ATOM 4878 CD2 LEU B 235 63.465 -6.607 78.795 1.00 27.4 ATOM 4878 CD2 LEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4889 CA ACH B 234 66.665 -9.191 78.462 1.00 27.4 ATOM 4889 CA CLEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4889 CD LEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA LEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA LEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA LEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA LEU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CA LEU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 27.4 ATOM 4880 CB LEU B 235 66.665 -9.191 78.664 1.00 27.4 ATOM 4880 CB LEU B 235 66.665 -9.191 78.462 1.00 27.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.662 1.00 27.4 ATOM 4880 CB LEU B 235 66.665 -9.191 78.662 1.00 27.4 ATOM 4880 CB LEU B 236 66.665 -9.191 78.662 1.00 27.4 ATOM 4880 CB LEU B 236 66.665 -9.191 78.662 1.00 27.4 ATOM 4880 CB LEU B 236 67.303 -10.206 77.629 1.00 24.4 ATOM 4889 CB LEU B 236 67.303 -10.206 77.629 1.00 24.4 ATOM 4889 CB LEYS B 237 67.449 -13.358 80.9	Mota								
ATOM 4860 CG PHE B 233	MOTA	4858	CA						
ATOM 4860 CG PHE B 233 71.217 -8.790 83.237 1.00 30.6 ATOM 4861 CD1 PHE B 233 72.285 -9.519 82.731 1.00 30.6 ATOM 4862 CD2 PHE B 233 70.481 -9.328 84.294 1.00 25.6 ATOM 4863 CE1 PHE B 233 70.481 -9.328 84.294 1.00 25.6 ATOM 4864 CE2 PHE B 233 70.481 -9.328 84.327 1.00 31.6 ATOM 4865 CE PHE B 233 70.803 -10.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 70.803 -10.573 84.832 1.00 31.6 ATOM 4866 C PHE B 233 71.871 -11.292 84.317 1.00 20.6 ATOM 4867 O PHE B 233 68.553 -9.567 81.270 1.00 21.6 ATOM 4868 N ALA B 234 65.535 -9.567 81.270 1.00 21.6 ATOM 4869 CA ALA B 234 65.855 -9.567 82.963 1.00 19.6 ATOM 4870 CB ALA B 234 65.835 -7.733 83.950 1.00 19.6 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.6 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 20.7 ATOM 4874 CA LEU B 235 66.822 -8.141 79.652 1.00 19.7 ATOM 4875 CB LEU B 235 66.822 -8.141 79.652 1.00 19.7 ATOM 4876 CG LEU B 235 63.465 -6.607 78.024 1.00 27.7 ATOM 4877 CDI LEU B 235 63.465 -6.607 78.024 1.00 27.7 ATOM 4878 CDI LEU B 235 63.465 -9.276 78.795 1.00 20.7 ATOM 4878 CD LEU B 235 63.465 -9.276 78.795 1.00 20.7 ATOM 4878 CD LEU B 235 63.465 -9.276 78.795 1.00 20.7 ATOM 4878 CD LEU B 235 63.783 -5.813 76.770 1.00 20.7 ATOM 4880 O LEU B 235 66.665 -9.917 78.462 1.00 27.7 ATOM 4881 N GLU B 236 66.665 -9.917 78.462 1.00 27.7 ATOM 4880 O LEU B 235 67.303 -10.206 77.629 1.00 27.7 ATOM 4881 N GLU B 236 66.665 -9.917 78.462 1.00 27.7 ATOM 4886 CG GLU B 236 67.303 -10.206 77.629 1.00 27.7 ATOM 4887 CC LEU B 235 67.404 88.00	MOTA	4859	CB						
ATOM		4860	CG	PHE I	B 233	71.217	-8.790		
ATOM 4862 CD2 PHE B 233				_		72.285	-9.519		1.00 30.48
ATOM 4863 CE1 PHE B 233						70.481	-9.328	84.294	1.00 25.32
ATOM 4864 CE2 PHE B 233		•						83.262	1.00 31.87
ATOM 4866 CZ PHE B 233									
ATOM 4866 C PHE B 233 68.712 -8.439 81.727 1.00 20.2 ATOM 4867 O PHE B 233 68.553 -9.567 81.270 1.00 21.2 ATOM 4868 N ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.4 ATOM 4872 O ALA B 234 65.772 -8.898 81.749 1.00 18.4 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 20.2 ATOM 4874 CA LEU B 235 66.634 -7.934 80.845 1.00 20.2 ATOM 4875 CB LEU B 235 64.822 -8.141 79.552 1.00 19.4 ATOM 4876 CG LEU B 235 63.465 -6.607 78.024 1.00 27.4 ATOM 4877 CD1 LEU B 235 63.465 -6.607 78.024 1.00 20.4 ATOM 4878 CD2 LEU B 235 62.761 -7.897 77.664 1.00 26.4 ATOM 4879 C LEU B 235 66.665 -9.191 78.462 1.00 19.4 ATOM 4880 O LEU B 235 66.665 -9.191 78.462 1.00 19.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4882 CA GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4888 CB GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4888 CB GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4889 O GLU B 236 68.969 -8.597 76.548 1.00 43.4 ATOM 4886 OE1 GLU B 236 68.969 -8.597 76.548 1.00 43.4 ATOM 4887 OE2 GLU B 236 66.665 -1.10 78.099 76.7551 1.00 22.4 ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 22.4 ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 22.4 ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 22.4 ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 23.4 ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 27.4 ATOM 4890 N LYS B 237 70.405 -13.683 80.924 1.00 31.4 ATOM 4891 CA LYS B 237 77.520 -11.748 79.492 1.00 24.4 ATOM 4893 CG LYS B 237 77.306 -12.806 80.993 1.00 23.4 ATOM 4896 NZ LYS B 237 77.520 -11.766 80.993 1.00 24.4 ATOM 4897 N SER B 238 62.776 -12.806 80.573 1.00 25.4 ATOM 4890 N SER B 238 62.776 -12.806 80.573 1.00 27.4 ATOM 4890 N SER B 238 62.776 -12.806 80.573 1.00 23.4 ATOM 4890 N SER B 238 62.776 -12.806 80.573 1.00 23.4 ATOM 4890 N SER B 238 62.776 -12.806 80.573 1.00 23.4 ATOM 4900 CA SER B 238 62.776 -12.806 80.573 1.00 23.4 ATOM 4900 CA SER B 238 62.776 -12.806 80.557 1.00 23.4 ATOM 4900 CA SER B 238 63.677 -13.860 80.6	MOTA	4864	CE2						
ATOM 4866 C PHE B 233 68.553 -9.567 81.270 1.00 21.5 ATOM 4868 N ALA B 234 67.842 -7.878 82.560 1.00 21.5 ATOM 4869 CA ALA B 234 66.626 -8.576 82.963 1.00 19.4 ATOM 4870 CB ALA B 234 65.835 -7.733 83.950 1.00 19.4 ATOM 4871 C ALA B 234 65.835 -7.733 83.950 1.00 19.4 ATOM 4871 C ALA B 234 65.772 -8.898 81.749 1.00 18.1 ATOM 4873 N LEU B 235 65.634 -7.934 80.845 1.00 20.3 ATOM 4873 N LEU B 235 64.822 -8.141 79.652 1.00 19.4 ATOM 4875 CB LEU B 235 64.773 -6.874 78.795 1.00 24.4 ATOM 4876 CG LEU B 235 64.773 -6.874 78.795 1.00 24.4 ATOM 4877 CD1 LEU B 235 63.465 -6.607 78.024 1.00 27.5 ATOM 4877 CD1 LEU B 235 62.761 -7.897 77.664 1.00 26.5 ATOM 4879 C LEU B 235 65.376 -9.276 78.795 1.00 26.5 ATOM 4880 O LEU B 235 66.665 -9.276 78.431 1.00 18.4 ATOM 4881 N GLU B 235 66.665 -9.276 78.431 1.00 18.5 ATOM 4882 CA GLU B 236 66.665 -9.191 78.462 1.00 19.5 ATOM 4883 CB GLU B 236 66.665 -9.191 78.462 1.00 19.5 ATOM 4886 CB GLU B 236 66.896 -8.597 76.548 1.00 31.5 ATOM 4888 C GLU B 236 68.969 -8.597 76.548 1.00 43.5 ATOM 4888 C GLU B 236 68.969 -8.597 76.558 1.00 48.7 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4889 C GLU B 236 67.2011.607 78.209 1.00 24.5 ATOM 4891 CA LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.607 79.763 1.00 29.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.00 24.5 ATOM 4894 CD LYS B 237 67.520 -11.748 79.492 1.	MOTA	4865	CZ	PHE 1					
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ATOM 4877 CD1 LEU B 235 63.783 -5.813 76.770 1.00 20. ATOM 4878 CD2 LEU B 235 62.761 -7.897 77.664 1.00 26. ATOM 4879 C LEU B 235 65.376 -9.276 78.795 1.00 20. ATOM 4880 O LEU B 235 64.648 -10.205 78.431 1.00 18. ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 19. ATOM 4882 CA GLU B 236 66.665 -9.191 78.462 1.00 19. ATOM 4883 CB GLU B 236 68.777 -9.853 77.384 1.00 31. ATOM 4884 CG GLU B 236 68.777 -9.853 77.384 1.00 31. ATOM 4885 CD GLU B 236 68.969 -8.597 76.548 1.00 43. ATOM 4886 OE1 GLU B 236 70.428 -8.292 76.259 1.00 45. ATOM 4887 OE2 GLU B 236 70.697 -7.309 75.538 1.00 48. ATOM 4888 C GLU B 236 67.2011.607 78.209 1.00 24. ATOM 4889 O GLU B 236 66.86£ -12.552 77.501 1.00 22. ATOM 4890 N LYS B 237 67.520 -11.748 79.492 1.00 24. ATOM 4891 CA LYS B 237 67.520 -11.748 79.492 1.00 24. ATOM 4892 CB LYS B 237 67.989 -12.984 81.562 1.00 23. ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4896 NZ LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4897 C LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4898 O LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4890 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4900 CA SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 CA SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 CA SER B 238 63.647 -13.280 79.229 1.00 28. ATOM 4900 CA SER B 238 63.677 -13.280 79.089 1.00 29. ATOM 4900 CA SER B 238 63.677 -13.280 79.089 1.00 29. ATOM 4900 CA SER B 238 63.677 -13.280 79.089 1.00 29. ATOM 4900 CA SER B 238 63.677 -13.280 79.089 1.00 29. ATOM 4900 CA SER B 238 63.677 -13.280 79.089 1.00 29. ATOM 4900 CA SER B 238 63.677 -13.280 79.089 1.00 29. ATOM 4900 CA SER B 238 63.				LEU	B 235	63.465	-6.607	78.024	1.00 27.87
ATOM 4877 CD2 LEU B 235 62.761 -7.897 77.664 1.00 26.4 ATOM 4879 C LEU B 235 65.376 -9.276 78.795 1.00 20.4 ATOM 4880 O LEU B 235 64.648 -10.205 78.431 1.00 18.4 ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4882 CA GLU B 236 66.665 -9.191 78.462 1.00 19.4 ATOM 4883 CB GLU B 236 68.777 -9.853 77.384 1.00 31.4 ATOM 4884 CG GLU B 236 68.777 -9.853 77.384 1.00 31.4 ATOM 4885 CD GLU B 236 68.969 -8.597 76.548 1.00 43.4 ATOM 4885 CD GLU B 236 70.697 -7.309 75.538 1.00 48.4 ATOM 4886 OE1 GLU B 236 70.697 -7.309 75.538 1.00 48.4 ATOM 4888 C GLU B 236 70.697 -7.309 75.538 1.00 48.4 ATOM 4888 C GLU B 236 67.2011.607 78.209 1.00 24.4 ATOM 4889 O GLU B 236 67.2011.607 78.209 1.00 24.4 ATOM 4891 CA LYS B 237 67.520 -11.748 79.492 1.00 24.4 ATOM 4891 CA LYS B 237 67.449 -13.058 80.130 1.00 27.4 ATOM 4893 CG LYS B 237 67.449 -13.058 80.130 1.00 27.4 ATOM 4894 CD LYS B 237 69.466 -12.641 81.650 1.00 29.4 ATOM 4894 CD LYS B 237 67.499 -12.984 81.562 1.00 23.4 ATOM 4896 NZ LYS B 237 70.305 -13.663 80.993 1.00 39.4 ATOM 4897 C LYS B 237 70.305 -13.663 80.993 1.00 39.4 ATOM 4898 O LYS B 237 70.305 -13.663 80.993 1.00 39.4 ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25.4 ATOM 4899 N SER B 238 63.677 -13.280 80.620 1.00 27.4 ATOM 4899 N SER B 238 63.677 -13.280 80.620 1.00 27.4 ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27.4 ATOM 4900 CA SER B 238 63.677 -12.806 80.565 1.00 29.4 ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23.4 ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23.4 ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23.4 ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23.4 ATOM 4900 CA SER B 238 62.776 -12.806 80.565 1.00 29.4 ATOM 4900 CA SER B 238 62.776 -12.806 79.763 1.00 27.4 ATOM 4900 CA SER B 238 62.776 -12.806 80.565 1.00 29.4 ATOM 4900 CA SER B 238 62.776 -12.806 79.089 1.00 23.4 ATOM 4900 CA SER B 238 62.387 79.13.192 75.836 1.00 27.4 ATOM 4900 CA SER B 238 62.776 -12.806 79.089 1.00 29.4 ATOM 4900 CA SER B 238 62.776 -12.806 79.089 1.00 29.4 ATOM 4900 CA SER B 238 62.75							-5.813	76.770	1.00 20.77
ATOM 4876 CD LEU B 235 65.376 -9.276 78.795 1.00 20. ATOM 4880 O LEU B 235 64.648 -10.205 78.431 1.00 18. ATOM 4881 N GLU B 236 66.665 -9.191 78.462 1.00 19. ATOM 4882 CA GLU B 236 67.303 -10.206 77.629 1.00 27. ATOM 4883 CB GLU B 236 68.777 -9.853 77.384 1.00 31. ATOM 4885 CD GLU B 236 68.969 -8.597 76.548 1.00 43. ATOM 4886 OE1 GLU B 236 70.428 -8.292 76.259 1.00 45. ATOM 4887 OE2 GLU B 236 70.697 -7.309 75.538 1.00 48. ATOM 4887 OE2 GLU B 236 70.697 -7.309 75.538 1.00 22. ATOM 4889 O GLU B 236 66.86£ -12.552 77.501 1.00 22. ATOM 4889 O GLU B 236 66.86£ -12.552 77.501 1.00 22. ATOM 4890 N LYS B 237 67.520 -11.748 79.492 1.00 24. ATOM 4891 CA LYS B 237 67.520 -11.748 79.492 1.00 24. ATOM 4892 CB LYS B 237 67.989 -12.984 81.562 1.00 23. ATOM 4894 CD LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4896 NZ LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4896 NZ LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4896 NZ LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4897 C LYS B 237 65.789 -14.766 79.763 1.00 39. ATOM 4899 N SER B 238 63.677 -12.806 80.573 1.00 25. ATOM 4899 N SER B 238 63.677 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4900 CA SER B 238 62.776 -12.241 81.289 1.00 29. ATOM 4900 C SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 C SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 C SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 C SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 C SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4900 C SER B 238 63.079 -13.192 75.886 1.00 32. ATOM 4906 CA LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 75.886 1.00 32.								77.664	1.00 26.94
ATOM 4887	MOTA								
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ATOM 4885 CD GLU B 236						68.969	-8.597	76.548	1.00 43.60
ATON 4886 OE1 GLU B 236							-8.292	76.259	1.00 45.19
ATOM 4887 OE2 GLU B 236 71.30C -9.032 76.751 1.00 52. ATOM 4888 C GLU B 236 67.2011.607 78.209 1.00 24. ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 22. ATOM 4890 N LYS B 237 67.520 -11.748 79.492 1.00 24. ATOM 4891 CA LYS B 237 67.449 -13.058 80.130 1.00 27. ATOM 4892 CB LYS B 237 67.989 -12.984 81.562 1.00 23. ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 63.677 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4902 OG SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4903 C SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.									1.00 48.77
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ATOM 4889 O GLU B 236 66.865 -12.552 77.501 1.00 22. ATOM 4890 N LYS B 237 67.520 -11.748 79.492 1.00 24. ATOM 4891 CA LYS B 237 67.449 -13.058 80.130 1.00 27. ATOM 4892 CB LYS B 237 67.989 -12.984 81.562 1.00 23. ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	ATOM	4888	С	GLU	B 236				
ATOM 4891 CA LYS B 237 67.520 -11.748 79.492 1.00 27. ATOM 4891 CA LYS B 237 67.449 -13.058 80.130 1.00 27. ATOM 4892 CB LYS B 237 67.989 -12.984 81.562 1.00 23. ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4903 C SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	MOTA	4889	0	GLU .	в 236				
ATOM 4891 CA LYS B 237 67.449 -13.058 80.130 1.00 27. ATOM 4892 CB LYS B 237 67.989 -12.984 81.562 1.00 23. ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 63.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.574 13.192 76.846 1.00 32.		4890	N	LYS	B 237			79.492	
ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4898 O LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 N SER B 238 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 63.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 75.837 1.00 30.						67.449	-13.058		1.00 27.10
ATOM 4893 CG LYS B 237 69.466 -12.641 81.650 1.00 29. ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 63.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.						67.989	-12.984	81.562	1.00 23.43
ATOM 4894 CD LYS B 237 70.305 -13.683 80.924 1.00 31. ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 75.837 1.00 30.								81.650	1.00 29.46
ATOM 4895 CE LYS B 237 71.782 -13.356 80.993 1.00 39. ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	MOTA					70.305	-13 693	80 924	1 00 31.65
ATOM 4895 NZ LYS B 237 72.580 -14.363 80.242 1.00 46. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	MOTA	4894	CD				-13.003		
ATOM 4896 NZ LYS B 237 72.580 -14.363 80.242 1.00 30. ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	MOTA	4895	CE	LYS	B 237	71.782			
ATOM 4897 C LYS B 237 66.019 -13.615 80.143 1.00 30. ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.776 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.		4896	NZ	LYS	B 237	72.580			1.00 46.74
ATOM 4898 O LYS B 237 65.789 -14.766 79.763 1.00 31. ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.						66.019	-13.615	80.143	1.00 30.92
ATOM 4899 N SER B 238 65.057 -12.806 80.573 1.00 25. ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.776 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.				TVC	B 237	65.789	-14.766	79.763	1.00 31.42
ATOM 4900 CA SER B 238 63.677 -13.280 80.620 1.00 27. ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4904 O SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	-			252	יכב ם				1.00 25.86
ATOM 4901 CB SER B 238 62.776 -12.241 81.289 1.00 23. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	MOTA			35K	2 22 2				1.00 27.98
ATOM 4901 CB SER B 238 62.776 -12.241 31.283 1.00 29. ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.	MOTA	4900	CA	SER	B 238	63.6//			1 00 23 89
ATOM 4902 OG SER B 238 62.756 -11.028 80.565 1.00 29. ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.		4901	CB						1.00 23.03
ATOM 4903 C SER B 238 63.145 -13.642 79.229 1.00 28. ATOM 4904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 4905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.			og			62.756			1.00 23.27
ATOM 1904 O SER B 238 62.387 -14.605 79.089 1.00 29. ATOM 1905 N LEU B 239 63.536 -12.886 78.203 1.00 27. ATOM 1906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.				SER	B 238	63.145	-13.642		1.00 28.32
ATOM 4904 O SER B 239 63.536 -12.886 78.203 1.00 27. ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32. ATOM 4906 CA LEU B 239 63.079 -13.192 75.837 1.00 30.				CEP	B 238		-14.605	79.089	1.00 29.65
ATOM 4906 CA LEU B 239 63.079 -13.192 76.846 1.00 32.				257	- 220			78.203	1.00 27.39
ATOM 9900 CA HEU B 222 (2 544 12 129 75 837 1 00 30.				1120	D 233		-13 102		1.00 32.52
ATOM 4907 CB LEU B 239 63.544 -12.129 /3.637 1.00 30.	MOTA	_		LEU	B 239		10 100		1 00 30.53
	MOTA	4907	CB	LEU	B 239	63.544	-12.129	, 5.657	1.00 30.32

ATOM	4908	CG LEU B 239	62.833 -10.772	75.895	1.00 36.06
ATOM	4909		63.404 -9.836	74.842	1.00 29.64
ATOM	4910				
			61.338 -10.976	75.667	1.00 30.73
MOTA	4911		63.598 -14.563	76.430	1.00 34.15
MOTA	4912		62.879 -15.340	75.803	1.00 33.23
ATOM	4913	N GLU B 240	64.844 -14.859	76.788	1.00 31.39
ATOM	4914	CA GLU B 240	65.434 -16.152	76,472	1.00 33.79
ATOM	4915		66.859 -16.238	77.011	1.00 38.51
ATOM	4916				
			67.878 -15.407	76.275	1.00 40.56
ATOM	4917		69.256 -15.532	76.903	1.00 48.20
ATOM	4918		69.685 -16.679	77.161	1.00 45.36
ATOM	4919	OE2 GLU B 240	69.912 -14.492	77.130	1.00 48.21
ATOM	4920	C GLU B 240	64.604 -17.258	77.108	1.00 33.23
MOTA	4921	O GLU B 240	64.391 -18.310	76.510	1.00 32.15
ATOM	4922	N ILE B 241	64.146 -17.017	78.331	1.00 29.72
ATOM	4923	CA ILE B 241	63.328 -17.989	79.047	1.00 29.85
	4924	CB ILE B 241			
ATOM			63.006 -17.489	80.466	1.00 30.42
ATOM	4925	CG2 ILE B 241	.62.049 -18.456	81.162	1.00 29.88
MOTA	4926	CG1 ILE B 241	64.309 -17.311	81.254	1.00 29.77
ATOM	4927	CD1 ILE B 241	64.118 -16.760	82.654	1.00 32.92
ATOM	4928	C ILE B 241	62.016 -18.247	78.298	1.00 34.65
MOTA	4929	O ILE B 241	61.592 -19.396	78.149	1.00 30.22
ATOM	4930	N VAL B 242	61.379 -17.178	77.823	1.00 31.16
ATOM	4931	CA VAL B 242	60.114 -17.312	77.105	1.00 34.55
ATOM	4932	CB VAL B 242	59.476 -15.937	76.825	
	4933	· - · - · - · - · - · - · - · · - ·			1.00 30.77
MOTA	-		58.191 -16.113	76.038	1.00 32.18
ATOM	4934	CG2 VAL B 242	59.201 -15.214	78.140	1.00 31.57
MOTA	4935	C VAL B 242	60.320 -18.042	75.787	1.00 36.56
MOTA	4936	O VAL B 242	59.572 -18.959	75.453	1.00 33,93
MOTA	4937	N LYS B 243	61.337 -17.627	75.042	1.00 38.64
MOTA	4938	CA LYS B 243	61.659 -18.241	73.760	1.00 44.36
ATOM	4939	CB LYS B 243	62.966 -17.659	73.214	1.00 48.33
ATOM	4940	CG LYS B 243	62.810 -16.399	72.386	1.00 53.88
	4941	CD LYS B 243	62.185 -16.718	71.036	
ATOM					1.00 53.72
ATOM	4942	CE LYS B 243	63.056 -17.681	70.242	1.00 54.69
ATOM	4943	NZ LYS B 243	62.456 -18.025	68.923	1.00 57.75
MOTA	4944	C LYS B 243	61.793 -19.755	73.824	1.00 43.21
MOTA	4945	O LYS B 243	61.432 -20.455	72.884	1.00 42.92
ATOM	4946	N GLU B 244	62.312 -20.257	74.935	1.00 45.77
ATOM	4947	CA GLU B 244	62.528 -21.687	75.085	1.00 47.72
ATOM	4948	CB GLU B 244	63.669 -21.925	76.075	1.00 50.89
ATOM	4949	CG GLU B 244	64.080 -23.378	76.208	1.00 57.16
	4950	CD GLU B 244	65.223 -23.564	77.173	1.00 57.74
MOTA					
ATOM	4951		66.295 -22.967	76.942	1.00 60.18
ATOM	4952	OE2 GLU B 244	65.049 -24.308	78.160	1.00 61.59
MOTA	4953	C GLU B 244	61.312 -22.507	75.505	1.00 47.78
ATOM	4954	O GLU B 244	61.376 -23.736	75.544	1.00 51.39
ATOM	4955	N VAL B 245	60.200 -21.851	75.805	1.00 43.31
ATOM	4956	CA VAL B 245	59.019 -22.589	76.230	1.00 43.55
ATOM	4957	CB VAL B 245	58.867 -22.514	77.771	1.00 45.89
ATOM	4958	CG1 VAL B 245	57.665 -23.322	78.231	1.00 49.90
	4959	CG2 VAL B 245	60.131 -23.040	78.435	1.00 46.37
ATOM				70.433	
ATOM	4960	C VAL B 245	57.727 -22.115	75.565	1.00 41.01
MOTA	4961	C VAL B 245	56.659 -22.676	75.798	1.00 39.36
MOTA	4962	N PHE B 246	57.814 -21.101	74.716	1.00 34.37
ATOM	4963	CA PHE B 246	56.610 ~20.602	74.077	1.00 34.36
ATOM	4964	CB PHE B 246	55.986 -19.517	74.958	1.00 30.80
MOTA	4965	CG PHE B 246	54.542 -19.230	74.644	1.00 32.57
ATOM	4966	CD1 PHE B 246	53.548 -20.142	74.989	1.00 27.72
	4967	CD2 PHE B 246	54.174 -18.048	74.003	1.00 28.20
ATOM			52.207 -19.878	74.704	
ATOM	4968				1.00 26.71
ATOM	1969	CE2 PHE B 246	52.836 -17.773	73.713	1.00 29.27
ATOM	4970	CZ PHE B 246	51.850 -18.689	74.065	1.00 26.18
· MOTA	4971	C PHE B 246	56.904 -20.040	72.682	1.00 35.32
MOTA	4972	O PHE B 246	57.740 -19.153	72.517	1.00 31.73
ATOM	4973	: GLU B 247	56.205 -20.568	71.683	1.00 37.00
				•	

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ATOM	497	4 C	A GLU	B 247	56.363	-20.137	70.296	1.00	40.73
MOTA	497	5 C	B GLU	B 247	56.518	-21.347	69.370		43.38
ATOM	497	6 C	G GLU	B 247		-22.702	70.073	1.00	51.49
ATOM	497	7 CI	D GLU	B 247		-23.214	70.718		55.29
MOTA	497	B 01	E1 GLU	B 247		-22.601	71.691		49.08
MOTA	497		E2 GLU			-24.246	70.241		60.95
ATOM	498		GLU	B 247		-19.379	69.939		39.04
ATOM	498:		GLU	B 247		-19.960	69.436		39.49
ATOM	4982	N S		B 248	55.076	-18.064	70.182		35.13
ATOM	4983	3 CI		B 248	56.188	-17.270	70.733		33.57
ATOM	4984	C.		B 248	53.935	-17.188	69.916		36.03
ATOM	4985	CE		B 248		-15.878	70.562		35.33
ATOM	4986	s co		B 248		-15.880	70.233		32.85
MOTA	4987	, c	PRO	B 248	53.563	-16.990	68.457		33.44
ATOM	4988	3 0	PRO	B 248		-16.808	67.604		29.38
ATOM	4989	N	GLU.	B 249	52.263	-17.012	68.182		32.23
MOTA	4990	CA	GLU	B 249	51.773	-16.782	66.828		29.35
ATOM	4991	CB	GLU :	B 249		-17.366	66.645		31.87
ATOM	4992	CG	GLU :	B 249		-18.867	66.787		28.64
ATOM	4993	CD	GLU :	B 249	48.847	-19.338	66.747		33.37
ATOM	4994	OE	1 GLU	B 249		-18.917	67.630		26.38
ATOM	4995	ΟE	2 GLU 1	B 249		-20.115	65.835		37.71
ATOM	4996	С	GLU 1	B 249		-15.273	66.650		28.25
ATOM	4997	0	GLU 1	B 249	51.776	-14.765	65.537	1.00	21.47
ATOM	4998	N	VAL I	3 250	51.561	-14.564	67.768	1.00	21.77
MOTA	4999	CA	VAL I	3 250		-13.110	67.756	1.00	21.41
ATOM	5000	CB	VAL I		50.027	-12.676	67.357	1.00	26.42
MOTA	5001	CG.	1 VAL E		49.037	-13.196	68.378	1.00	20.96
MOTA	5002	CG:			49.931		67.243	1.00	24.68
MOTA	5003	С	VAL E			-12.608	69.168	1.00	22.88
MOTA	5004	0	VAL -E		51.592	-13.354	70.133	1.00	18.00
MOTA	5005	N	TYR E			-11.359	69.295		20.28
MOTA	5006	CA	TYR E			-10.823	70.620		22.33
MOTA	5007	ĊВ	TYR E			-11.043	70.999		20.67
ATOM	5008	CG				-10.045	70.427		21.89
ATOM	5009	CD			55.198	-8.834	71.072		19.31
ATOM	5010	CEI			56.129	-7.922	70.562		23.73
ATOM	5011	CD2				-10.321	69.254		18.72
MOTA	5012	CE2			56.580	-9.417	68.734		22.71
ATOM	5013	CZ	TYR B		56.813	-8.220	69.390		27.33
ATOM	5014	ОН	TYR B		57.705	-7.308	68.865		23.18
MOTA	5015 5016	С 0	TYR B		52.134 52.095	-9.349	70.732 69.728		25.71
MOTA MOTA	5017	N	LEU B		51.834	-8.622 -8.930	71.958		20.14
ATOM	5018	CA	LEU B		51.533	-7.532	72.252		21.13
ATOM	5019	CB	LEU B		50.154	-7.373	73.897		22.88
ATOM	5020	CG	LEU B		48.915	-7.435	1.996		23.73
ATOM	5021	CD1		252	48.779	-8.792	71.360		23.18
ATOM	5022	CD2		252	47.697	-7.119	72.833		29.06
ATOM	5023	c	LEU B	252	52.610	-7.044	73.217		24.77
ATOM	5024	ō	LEU B		53.064	-7.797	74.076		23.33
ATOM	5025	N	LEU B		53.011	-5.786	73.071		20.14
MOTA	5026	CA	LEU B		54.057	-5.209	73.911		20.33
MOTA	5027	CB	LEU B		55.304	-4.946 ·	73.051		15.18
MOTA	5028	CG	LEU B		56.490	-4.210	73.688		18.34
ATOM	5029		LEU B		57.062	-5.044	74.829		14.11
MOTA	5030		LEU B		57.552	-3.953	72.624	1.00	
MOTA	5031	c	LEU B		53.550	-3.913	74.536	1.00	
ATOM	5032	ō	LEU B		53.200	-2.974	73.821	1.00	
ATOM	5033	N	GLN B		53.495	-3.858	75.865	1.00	
ATOM	5034	CA	GLN B		53.000	-2.654	76.539		21.77
ATOM	5035	CB	GLN B		52.129	-3.040	77.755	1.00	
ATOM	5036	CG	GLN B		52.724	-2.815	79.124		32.51
ATOM ·	5037	CD	GLN B		52.563	-1.396	79.609		28.19
ATOM	5038	OE1	GLN B		51.507	-0.996	80.124		26.96
MOTA	5039	NE2	GLN B	254	53.603	-0.619	79.432	1.00	

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WO 01/018045

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1.00 20.15
                                            -1.793
                                                     76.887
                                   54.211
                   GLN B 254
MOTA
        5040
              С
                                   55.186
                                                     77.497
                                                              1.00 20.11
                                            -2.254
                   GLN B 254
        5041
              0
MOTA
                                                              1.00 19.46
                                            -0.532
                                                     76.468
                  LEU B 255
                                   54.146
        5042
              N
MOTA
                                             0.386
                                                     76.614
                                                              1.00 15.99
                   LEU B 255
                                   55.268
MOTA
        5043
              CA
                                                              1.00 18.15
                                   55.692
                                             0.831
                                                     75.211
                   LEU B 255
        5044
              CB
ATOM
                                                     74.296
                                                              1.00 21.80
                   LEU B 255
                                   56.143
                                            -0.316
        5045
              CG
ATOM
                                                     72.850
                                                              1.00 16.70
                                             0.159
                  LEU B 255
                                   56.215
       5046
              CD1
MCTA
                                                     74.771
                                                              1.00 13.76
                                            -0.843
                  LEU B 255
                                   57.501
        5047
              CD2
MOTA
                                                              1.00 21.41
                                                     77.492
                   LEU B 255
                                   55.083
                                             1.614
        5048
              C
ATOM
                                   55.379
                                             2.741
                                                     77.065
                                                              1.00 18.40
                   LEU B 255
        5049
              0
ATCM
                                                     78.718
                                                              1.00 16.80
                                             1.408
                                   54.618
        5050
              N
                   GLY B 256
ATOM
                                                              1.00 19.90
                                                     79.634
                   GLY B 256
                                   54.456
                                             2.519
              CA
MOTA
        5051
                                                              1.00 17.68
                   GLY B 256
                                             3.181
                                                     79.818
                                   55.816
        5052
              C
ATOM
                                                              1.00 13.96
                                   56.854
                                             2.514
                                                     79.841
                   GLY B 256
        5053
              0
ATOM
                                             4.497
                                                     79.936
                                                              1.00 19.55
                   THR B 257
                                   55.824
        5054
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MOTA
                                                     80.098
                                                              1.00 19.47
                   THR B 257
                                   57.081
                                             5.205
        5055
              CA
MOTA
                                                     79.340
                                                              1.00 21.49
                                   57.044
                                             6.547
                   THR B 257
        5056
              CB
ATOM
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                                                              1.00 17.43
                                   55.989
                                             7.365
        5057
              OG1
                   THR B 257
MOTA
                                                              1.00 22.49
                                                     77.850
                                   56.780
                                             6.311
                  THR B 257
ATOM
        5058
              CG2
                                                              1.00 20.75
                                             5.466
                                                     81.564
                   THR B 257
                                   57.440
        5059
ATOM
                                                              1.00 25.01
                                   58.480
                                             6.054
                                                     81.843
                   THR B 257
        5060
              0
MOTA
                                                     82.504
                                                              1.00 17.23
                                             5.004
                   ASP B 258
                                   56.618
        5061
              Ν
ATOM
                                                              1.00 17.42
                   ASP B 258
                                             5.277
                                                     83.906
                                   56.929
        5062
              CA
ATOM
                                             4.940
                                                     84.846
                                                              1.00 12.75
                         258
                                   55.744
              CB
                   ASP B
        5063
ATOM
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                                                     84.676
                                                              1.00 21.60
                                   55.197
                   ASP B 258
        5064
              ÇG
ATOM
                                                              1.00 16.09
                                                     84.460
                                             4.718
                   ASP B 258
                                    58.245
        5065
              C
MOTA
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                                                     85.542
                   ASP B 258
                                    58.667
                                             5.116
        5066
              0
ATOM
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                                             2.642
                                                     84.150
                                                               1.00 17.74
                  ASP B 258
ATOM
        5067
              OD1
                                                               1.00 18.68
                                                     85.109
                                             3.281
                   ASP B 258
                                    54.041
        5068
              OD2
ATOM
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                                                     83.779
                   PRO B 259
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        5069
              N
ATOM
                                                               1.00 17.75
                                    58.474
                                             2.901
                                                     82.641
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              CD
MOTA
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                                                     84.321
                   PRO B 259
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        5071
              CA
ATOM
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                                                               1.00 23.46
                                             1.988
                   PRO B 259
                                    60.395
              CB
        5072
ATOM
                                                               1.00 27.08
                                    59.800
                                             2:343
                                                     82.199
                   PRO B 259
        5073
              CG
ATOM
                                    61.305
                                                     84.172
                                                               1.00 23.86
                                             4.284
        5074
              С
                   PRO B 259
ATOM
                                             4.082
                                                               1.00 24.24
                                                     84.698
                                    62.406
        5075
              0
                   PRO B 259
MOTA
                                                               1.00 20.49
                                                     83.465
                   LEU B 260
                                    61.054
                                             5.387
        5076
              N
ATOM
                                    62.080
                                             6.417
                                                     83.262
                                                               1.00 15.17
                         260
        5077
              CA
                   LEU B
MOTA
                                    61.626
                                             7.408
                                                     82.185
                                                               1.00 17.03
        5078
              CB
                   LEU B 260
ATOM
                                                               1.00 16.02
                                             6.881
                                                     80.760
                   LEU B 260
                                    61.431
        5079
              CG
ATOM
                                                     79.901
                                                               1.00 17.03
                   LEU B 260
                                    60.703
                                             7.915
              CDI
        5080
ATOM
                                                               1.00 18.58
                                    62.803
                                              6.546
                                                     80.163
        5081
              CD2
                   LEU B 260
ATOM
                                    62.449
                                                     84.541
                                                               1.00 22.45
                                              7.194
                   LEU B
                         260
        5082
ATOM
                                             7.440
                                                     85.412
                                                               1.00 17.84
                   LEU B 260
                                    61.611
              0
        5083
ATOM
                                                     84.635
                                                               1.00 22.90
                                    63.713
                                              7.588
                   LEU B 261
        5084
              N
ATOM
                                              8.332
                                                     85.782
                                                               1.00 26.34
                                    64.219
        3085
              CA
                   LEU B
                         261
ATOM
                                                     85.473
                                                               1.00 20.58
                                             8.914
                                    65.605
                   LEU B 261
        5086
              CB
ATOM
                                                     86.553
                                             9.850
                                                               1.00 28.44
                                    66.180
                   LEU B 261
        3087
              CG
ATOM
                                    66.481
                                                               1.00 29.84
                                              9.055
                                                     87.812
                         261
        5088
              CD1
                  LEU B
ATOM
                                    67.462
                                            10.522
                                                     86.057
                                                               1.00 32.10
                   LEU B 261
        5089
              CD2
ATOM
                                             9.475
                                                     86.227
                                                               1.00 27.61
                                    63.315
              C
                   LEU B 261
ATOM
        5090
                                                               1.00 24.02
                                                     87.408
                                    62.978
                                             9.586
                         261
        5091
              0
                   LEU B
MOTA
                                                     85.269
                                                               1.00 23.33
                                    62.934
                                            10.315
                   GLU B 262
        5092
ATOM
                                            11.490
                                                     85.530
                                                               1.00 23.38
                                    62.126
              CA
                   GLU B 262
        5093
ATOM
                                                     84.302
                                                               1.00 23.17
                                    62.115
                                            12.415
                   GLU B
                         262
        5094
              CB
MOTA
                                                     83.806
                                                               1.00 28.98
                                    63.503
                                            12.854
        5095
              CG
                   GLU B 262
ATOM
                                                     82.902
                                                               1.00 32.26
                                            11.831
                                    64.179
              CD
                   GLU B 262
        5096
ATOM
                                                               1.00 29.28
                                                      82.838
                                    63.702
                                            10.673
                   GLU B
                         262
        5097
              CE1
ATOM
                                                     82.264
                                                               1.00 25.42
                                    65.201
                                            12.186
              OE2
                   GLU B 262
        5098
ATOM
                                                               1.00 23.25
                                            11.249
                                                      85.976
                                    60.693
              C
                   GLU B 262
        5099
STOM
                                                               1.00 27.63
                                                      86.368
                                    60.013
                                            12.192
                   GLU B
                         262
        5100
              0
ATOM
                                                      85.927
                                                               1.00 22.25
                                    60.219
                                            10.011
              N
                   ASP B 263
        5101
ATOM
                                             9.751
                                                      86.345
                                                               1.00 24.46
                                    58.840
              CA
                   ASP
                       В
                         263
        5102
ATOM
                                                               1.00 20.94
                                                      85.465
                                              8.659
                                    58.214
                   ASP B
              CB
                         263
        5103
ATOM.
                                                               1.00 25.30
                                    56.710
                                              8.543
                                                      85.659
              CG
                   ASP B 263
        5104
ATOM
                                              8.318
                                                     84.656
                                                               1.00 21.82
                                    55.995
                  ASP
                         263
ATOM
        5105
              OD1
                       В
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Figure 18-78

MOTA	5106	OD2 ASP B 26	3 56.239	8.666	86.811	1.00 18.31
MOTA	5107			9.339	87.814	1.00 26.39
MOTA	5108			8.335	88.179	1.00 22.11
ATOM ATOM	5109 5110	_		10.124 9.864	88.648 90.084	1.00 25.81
MOTA	5111	CB TYR B 26		11.055	90.841	1.00 36.80
MOTA	5112	CG TYR B 26		12.356	90.645	1.00 46.58
MOTA	5113	CD1 TYR B 26		13.166	89.542	1.00 47.03
MOTA	5114	CE1 TYR B 26		14.370	89.364	1.00 50.25
MOTA	5115	CD2 TYR B 26 CE2 TYR B 26		12.779	91.565	1.00 50.94
MOTA MOTA	5116 5117	CE2 TYR B 26 CZ TYR B 26		13.977 14.769	91.396 90.297	1.00 51.28 1.00 52.21
ATOM	5118	OH TYR B 26		15.961	90.142	1.00 49.65
ATOM	5119	C TYR B 26	4 57.340	8.628		1.00 31.04
MOTA	5120	O TYR B 26		8.181	91.657	1.00 24.50
ATOM	5121	N LEU B 26		8.074	89.666	1.00 26.68
ATOM ATOM	5122 5123	CA LEU B 26 CB LEU B 26		6.900 6.838	90.086 89.390	1.00 24.17 1.00 24.69
ATOM	5124	CG LEU B 26		7.982	89.761	1.00 26.00
MOTA	5125	CD1 LEU B 26		7.583	89.460	1.00 22.21
MOTA	5126	CD2 LEU B 26		8.281	91.238	1.00 29.31
MOTA	5127	C LEU B 26		5.568	89.948	1.00 25.83
MOTA MOTA	5128 5129	O LEU B 269 N SER B 269		4.512 5.618	89.908 89.867	1.00 21.74 1.00 23.30
ATOM	5130	CA SER B 26		4.398	89.813	1.00 20.75
ATOM	5131	CB SER B 26	58.820	3.900	88.378	1.00 19.67
MOTA	5132	OG SER B 26		4.615	87.739	1.00 18.11
MOTA	5133	C SER B 266		4.710	90.420	1.00 23.01
ATOM ATOM	5134 5135	O SER B 266 N LYS B 267		5.845 3.707	90.345 91.023	1.00 17.74 1.00 24.25
ATOM	5136	CA LYS B 267		3.916	91.613	1.00 23.79
ATOM	5137	CB LYS B 267		3.153	92.929	1.00 23.71
MOTA	5138	CG LYS B 267		3.582	93.960	1.00 27.29
ATOM	5139 5140	CD LYS B 267 CE LYS B 267		5.088 5.535	94.207 95.273	1.00 30.33
MOTA MOTA	5141	NZ LYS B 267		7.004	95.509	1.00 33.37
ATOM	5142	C LYS B 267		3.483	90.634	1.00 26.41
ATOM	5143	O LYS B 267		3.317	91.016	1.00 25.33
ATOM	5144	N PHE B 268		3.288	89.375	1.00 22.18
MOTA MOTA	5145 5146	CA PHE B 268		2.919 2.171	88.318 87.179	1.00 22.78 1.00 20.55
ATOM	5147	CG PHE B 268		0.761	87.526	1.00 19.23
ATOM	5148	CD1 PHE B 268	61.722	-0.025	86.585	1.00 20.72
MOTA	5149	CD2 PHE B 268		0.207	88.773	1.00 18.17
ATOM	5150	CE1 PHE B 268 CE2 PHE B 268		-1.336 -1.105	86.875 89.073	1.00 18.83
ATOM ATOM	5151 5152	CE2 PHE B 268		-1.105	88.122	1.00 20.05
ATOM	5153	C PHE B 268		4.222	87.785	1.01 23.66
ATOM	5154	O PHE B 268		5.232	87.692	1.00 19.40
ATOM	5155	N ASN B 269	. 65.396	4.203	87.437	1.00 21.96
MOTA	5156	CA ASN B 269		5.396	86.926 87.824	1.00 25.04 1.00 25.68
ATOM ATOM	5157 5158	CB ASN B 269 CG ASN B 269	67.243 66.845	5.783 5.946	89.273	1.00 25.08
ATOM	5159	OD1 ASN B 269		6.557	89.579	1.00 28.81
ATOM	5160	ND2 ASN B 269	6.7.659	5.419	90.176	1.00 31.12
MOTA	5161	C ASN B 269	66.579	5.151	85.523	1.00 25.87
ATOM	5162	O ASN B 269	67.769	5.336	85.268	1.00 24.58
MOTA	5163 5164	N LEU B 270 CA LEU B 270	65.695 66.116	4.757 4.462	84.611 83.241	1.00 21.37 1.00 16.35
MOTA MOTA	5165	CB LEU B 270	65.176	3.426	82.610	1.00 24.12
ATOM	5166	CG LEU B 270	64.909	2.144	83.412	1.00 27.89
ATOM	5167	CDI LEU B 270	64.181	1.136	82.515	1.00 23.01
ATOM	5168	CD2 LEU B 270	66.221 66.184	1.547	83.904 82.337	1.00 23.92 1.00 20.06
MOTA	5169 5170	C LEU B 270 O LEU B 270	65.654	5.682 6.761	82.663	1.00 20.06
ATOM ATOM	5170	N SER B 271	66.839	5.497	81.193	1.00 20.07
A1 041					•	

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			onn n 271	66.989	6.546	80,200	1.00 21.20
MOTA	5172	CA	SER B 271 SER B 271	68.437	6.621	79,714	1.00 21.80
MOTA	5173	CB		68.772	5.485	78.921	1.00 21.47
MOTA	5174	OG		66.106	6.228	79.000	1.00 22.83
ATOM	5175	C	SER B 271	65.631	5.102	78.854	1.00 16.12
MOTA	5176	0	SER B 271	65.916	7.238	78.154	1.00 20.84
MOTA	5177	N	ASN B 272	65.152	7.156	76.906	1.00 27.82
ATOM	5178	CA	ASN B 272		8.478	76.123	1.00 30.30
MOTA	5179	CB	ASN B 272	65.263	9.456	76.475	1.00 37.83
MOTA	5180	CG.	ASN B 272	64.198	10.575	75.946	1.00 37.72
ATOM	5181	OD1	ASN B 272	64.167	9.052	77.360	1.00 41.69
ATOM	5182	ND2	ASN B 272	63.299	6.088	75.974	1.00 26.88
ATOM	5183	С	ASN B 272	65.701	5.280	75.412	1.00 23.12
MOTA	5184	0	ASN B 272	64.967		75.774	1.00 20.40
MOTA	5185	N	VAL B 273	67.012	6.160	74.899	1.00 27.34
ATOM	5186	CA	VAL B 273	67.745	5.260	74.805	1.00 30.40
ATOM	5187	CB	VAL B 273	69.225	5.705	74.029	1.00 34.98
ATOM	5188	CG1	VAL B 273	70.036	4.691	74.115	1.00 33.57
MOTA	5189	CG2	VAL B 273	69.299	7.057	75.343	1.00 24.23
ATOM	5190	С	VAL B 273	67.664	3.812	74.513	1.00 24.19
ATOM	5191	0	VAL B 273	67.590	2.913	76.648	1.00 20.96
ATOM	5192	N	ALA B 274	67.690	3.580	77.151	1.00 18.12
MOTA	5193	CA	ALA B 274	67.589	2.220		1.00 19.09
MOTA	5194	CB	ALA B 274	67.858	2.195	78.646	1.00 18.23
ATOM	5195	С	ALA B 274	66.172	1.729	76.863	1.00 20.77
ATOM	5196	0	ALA B 274	65.962	0.567	76.525	1.00 18.50
ATOM	5197	N	PHE B 275	65.207	2.631	77.003	1.00 21.25
ATOM	5198	CA	PHE B 275	63.802	2.310	76.761	1.00 22.24
ATOM	5199	CB	PHE B 275	62.941	3.546	77.037	1.00 24.72
ATOM	5200	CG	PHE B 275	61.466	3.303	76.921	1.00 23.64
ATOM	5201	CD1	PHE B 275	60.815	2.483	77.826	1.00 27.31
MOTA	5202	CD2	PHE B 275	60.732	3.893	75.907	1.00 27.82
ATOM	5203	CE1	PHE B 275	59.450	2.254	77.722	1.00 27.62
ATOM	5204	CE2		59.365	3.670	75.795	1.00 27.02
MOTA	5205	CZ	PHE B 275	58.72 7	2.851	76.701	1.00 24.47
ATOM	5206	Ċ	PHE .B 275	63.642	1.860	75.305	1.00 22.68
MOTA	5207	0	PHE B 275	63.045	0.821	75.030 74.378	1.00 23.85
ATOM	5208	N	LEU B 276	64.183	2.648	_	1.00 21.28
ATOM	5209	CA	LEU B 276	64.128	2.330	72.946 72.134	1.00 19.87
MOTA	5210	CB	LEU B 276	64.814	3.421	70.662	1.00 24.94
ATOM	5211	CG	LEU B 276	65.114	3.132	69.936	1.00 24.81
ATOM	5212	CD1	LEU B 276	63.818	2.852	70.018	1.00 21.01
MOTA	5213	CD2	LEU B 276	65.840	4.312	72.653	1.00 22.33
MOTA	5214	C	LEU B 276	64.841	1.021	71.886	1.00 20.73
ATOM	5215	0	LEU B 276	64.348	0.191	73.261	1.00 20.72
ATOM	5216	N	LYS B 277	66.011	0.857	73.201	1.00 24.36
ATOM	5217	CA	LYS 3 277	66.823	-0.335	73.938	1.00 27.37
ATOM	5218	CB	LYS B 277	68.086	-0.239	73.3381	1.00 35.58
MOTA	5219	CG	LYS 3 277	69.303	-0.973		
ATOM	5220	CD	LYS B 277	69.061	-2.456	73.188	
ATOM	5221	CE	LYS B 277	70.283	-3.137	72.580 71.230	
ATOM	5222	NZ	LYS B 277	70.616	-2.586		
MOTA	5223	С	LYS B 277	66.000	-1.554	73.482	00
	5224	0	LYS B 277	65.987	-2.568	72.777	
MOTA	5225	N	ALA B 278	65.319	-1.454		
ATOM	5226	CA	ALA B 278	64.476	-2.544		
MOTA	5227	СВ	ALA B 278	63.752	-2.117		
MOTA	5228	c	ALA B 278	63.459	-2.896	74.031	
MOTA	5229	ŏ	ALA B 278	63.231	-4.068		
MOTA	5230	N	PHE B 279	62.849	-1.862		74
ATOM	5231	CA	PHE B 279	61.860	-2.014		
ATOM		CB	PHE B 279	61.395	-0.629		
ATOM	5232 5233	CG	PHE B 279	60.467	-0.640		
ATOM		CD.	1 PHE B 279	59.196	-1.182	70.882	
ATOM	5234	כם:	2 PHE B 279	60.862	-0.078	69.567	
MOTA	5235	رت. دي	1 PHE B 279	58.325	-1.162	69.799	
ATOM	5236	CE.	2 PHE B 279	60.001	-0.051	68.476	1.00 25.57
ATOM	5237	.ندب				-	

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ATOM 5238 CZ PHE B 279 58.727 -0.594 68.592 1.00 25.13 ATOM 5240 O PHE B 279 61.866 -3.697 70.678 1.00 23.65 ATOM 5241 N ASN B 280 63.677 -2.376 70.804 1.00 22.65 ATOM 5242 CA ASN B 280 65.520 -2.306 69.660 1.00 22.37 ATOM 5244 CQ ASN B 280 65.520 -2.248 69.164 1.00 22.61 ATOM 5246 ND2 ASN B 280 64.746 -1.466 7.756 7.796 1.00 26.54 ATOM 5248 N 1LE B 281 65.408 -4.724 67.714 1.00 26.16 ATOM 5250 CA ILE B 281 65.408 -6.007 71.167 1.00 26.16 ATOM 5251 CB ILE B 281 66.006 -7.527 73.648 1.00 28								eo - co	1 00 05 13
ATOM 5239 C PHE B 279 62.472 -2.768 71.212 1.00 24.50 ATOM 5241 N ASN B 280 63.677 -2.376 70.878 1.00 24.54 ATOM 5242 CA ASN B 280 64.318 -3.046 69.680 1.00 21.93 ATOM 5243 CB ASN B 280 65.520 -2.248 69.680 1.00 23.73 ATOM 5243 CB ASN B 280 65.520 -2.248 69.680 1.00 23.73 ATOM 5244 CQ ASN B 280 65.520 -2.248 69.680 1.00 23.73 ATOM 5245 OD1 ASN B 280 65.520 -0.0376 68.505 1.00 25.81 ATOM 5246 ND2 ASN B 280 65.507 -0.937 68.505 1.00 25.81 ATOM 5246 ND2 ASN B 280 64.746 -4.466 70.009 1.00 26.10 ATOM 5249 N ILE B 281 65.485 -6.657 71.667 1.00 26.54 ATOM 5249 N ILE B 281 65.485 -6.657 71.667 1.00 25.81 ATOM 5250 CA ILE B 281 65.485 -6.607 71.667 1.00 25.81 ATOM 5251 CB ILE B 281 66.046 -7.527 73.648 1.00 28.50 ATOM 5253 CGI ILE B 281 66.046 -7.527 73.648 1.00 28.50 ATOM 5255 C ILE B 281 66.046 -7.527 73.648 1.00 28.50 ATOM 5255 C ILE B 281 64.320 -7.100 71.507 1.00 25.77 ATOM 5255 C ILE B 281 64.320 -7.100 71.507 1.00 25.77 ATOM 5255 C ILE B 281 64.320 -7.100 71.507 1.00 25.77 ATOM 5255 C ILE B 281 64.484 -8.11 70.992 1.00 23.07 ATOM 5255 C ILE B 282 61.4320 -7.700 71.507 1.00 25.77 ATOM 5255 C ILE B 282 61.4320 -7.700 71.507 1.00 25.77 ATOM 5255 C ILE B 282 61.44 320 -7.700 71.507 1.00 25.77 ATOM 5255 C ILE B 282 61.44 320 -7.700 71.507 1.00 25.77 ATOM 5256 CGI VAL B 282 60.703 -6.687 71.990 1.00 24.07 ATOM 5257 C VAL B 282 60.703 -7.700 71.507 1.00 25.77 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 25.77 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 24.07 ATOM 5257 C VAL B 282 60.703 -7.700 71.507 1.00 25.77 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 24.07 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 24.07 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 24.07 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 24.07 ATOM 5260 CGI VAL B 282 60.703 -7.700 71.507 1.00 25.77 ATOM 5260 CGI VAL B 282 60.703 -7.700 60.700 1.00 25.70 ATOM 5270 CG VAL B 282 60.700 -7.700 71.507 1.00 25.68 ATOM 5260 CGI VAL B 282 60.700 -7.700 60.700 1.00 25.70 ATOM 5270 CG VAL B 282 60.700 60.700 60.700 60.700 60.700 6	ATOM	5238	CZ	PHE B	279	58.727			
ATOM 5240 O PHE B 279 61.866 -3.697 70.678 1.00 26.54 ATOM 5241 N ASN B 280 63.677 -2.376 70.804 1.00 21.93 ATOM 5242 CA ASN B 280 65.520 -2.248 69.164 1.00 22.63 ATOM 5243 CB ASN B 280 65.520 -2.248 69.164 1.00 22.63 ATOM 5245 ODI ASN B 280 65.520 -2.248 69.164 1.00 22.63 ATOM 5246 ND2 ASN B 280 65.507 -0.937 68.505 1.00 30.83 ATOM 5246 ND2 ASN B 280 65.507 -0.937 68.505 1.00 30.83 ATOM 5246 ND2 ASN B 280 65.500 -0.112 68.714 1.00 26.54 ATOM 5247 C ASN B 280 64.775 -5.321 69.124 1.00 26.54 ATOM 5248 O ASN B 280 64.775 -5.321 69.124 1.00 26.16 ATOM 5248 O ASN B 280 64.775 -5.321 69.124 1.00 26.16 ATOM 5250 CA ILLE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5250 CA ILLE B 281 65.080 -4.724 71.272 1.00 26.16 ATOM 5255 CG ILLE B 281 66.086 -6.099 73.124 1.00 28.53 ATOM 5252 CGZ ILLE B 281 66.086 -6.099 73.124 1.00 28.53 ATOM 5255 C ILLE B 281 66.086 -7.527 73.648 1.00 28.53 ATOM 5255 C ILLE B 281 66.086 -7.527 73.648 1.00 28.53 ATOM 5255 C ILLE B 281 66.086 -7.527 73.678 1.00 23.077 ATOM 5255 C ILLE B 281 64.484 -7.030 71.507 1.00 25.77 ATOM 5256 O ILLE B 281 64.484 -7.030 71.507 1.00 25.77 ATOM 5256 C ILLE B 281 64.484 -7.030 71.507 1.00 23.07 ATOM 5256 C ILLE B 282 61.961 -7.465 71.813 1.00 22.04 ATOM 5256 C ILLE B 282 61.961 -7.465 71.813 1.00 22.04 ATOM 5256 C ILLE B 282 61.961 -7.465 71.813 1.00 22.05 ATOM 5261 CGZ VAL B 282 61.961 -7.465 71.813 1.00 22.06 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.06 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.06 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.06 ATOM 5263 O VAL B 282 61.961 -7.465 71.813 1.00 22.06 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.06 ATOM 5262 C C ALB 68.83 61.576 -6.971 68.660 1.00 27.95 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.65 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.65 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.65 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.65 ATOM 5262 C VAL B 283 61.960 -7.775 72.367 1.00 23.80 60.93 1.00 23.80 60.93 1.00 23.80 60.93 1.00 23.80 60.93 1.00 23.80 6	MOTA	5239	С	PHE B	279	62.472	-2.768	71.212	
ATOM 5241 CA ASN B 280 64.318 -3.046 69.680 1.00 21.93 ATOM 5243 CB ASN B 280 65.520 -2.248 69.644 1.00 22.63 ATOM 5244 CQ ASN B 280 65.520 -2.248 69.164 1.00 23.70 ATOM 5245 ODI ASN B 280 65.520 -2.248 69.164 1.00 23.63 ATOM 5246 ND2 ASN B 280 65.520 -0.379 68.505 1.00 25.81 ATOM 5247 C ASN B 280 64.746 -4.466 70.009 1.00 26.10 ATOM 5248 O ASN B 280 64.746 -4.466 70.009 1.00 26.10 ATOM 5249 N ILE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5249 N ILE B 281 65.080 -4.724 71.272 1.00 26.15 ATOM 5250 CA ILE B 281 66.006 -6.087 73.124 1.00 28.53 ATOM 5251 CB ILE B 281 66.006 -6.087 73.124 1.00 28.53 ATOM 5252 CG2 ILE B 281 66.006 -6.087 73.124 1.00 28.53 ATOM 5252 CG2 ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 66.036 -7.527 73.648 1.00 28.24 ATOM 5255 C ILE B 281 66.385 -7.427 74.541 1.00 28.53 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 25.73 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 25.73 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 25.73 ATOM 5256 C ILE B 281 64.320 -7.030 71.507 1.00 25.73 ATOM 5258 CA VAL B 282 63.139 -6.618 71.950 1.00 23.39 ATOM 5258 CA VAL B 282 63.139 -6.618 71.950 1.00 23.39 ATOM 5258 C VAL B 282 61.961 -7.465 71.813 1.00 22.90 ATOM 5260 CCI VAL B 282 60.703 -6.775 70.399 1.00 23.87 ATOM 5260 CCI VAL B 282 60.676 -6.577 79.594 88 1.00 23.87 ATOM 5261 C VAL B 282 60.676 -6.577 90.339 1.00 22.28 ATOM 5262 C VAL B 282 60.676 -6.577 90.339 1.00 22.28 ATOM 5266 C C ARB 8 283 60.576 -6.697 73.906 1.00 24.37 ATOM 5266 C C ARB 8 283 60.576 -6.697 73.906 1.00 26.45 ATOM 5267 C C ARB 8 283 60.576 -6.697 73.906 1.00 26.45 ATOM 5268 C C ARB 8 283 60.576 -6.697 73.906 1.00 26.45 ATOM 5266 C C ARB 8 283 60.576 -6.587 73.906 1.00 26.45 ATOM 5267 C C ARB 8 283 60.576 -6.597 73.906 1.00 26.45 ATOM 5267 C C ARB 8 283 60.501 -7.795 69.948 1.00 22.68 ATOM 5268 C C ARB 8 283 60.501 -7.795 69.948 1.00 22.68 ATOM 5267 C C ARB 8 283 60.501 -7.795 69.978 1.00 25.48 ATOM 5268 C C ARB 8 283 60.501 -7.795 69.998 1.00 25.48 ATOM 5267 C C ARB 8 283 60.501 -7.795 69.998 1.00 25.48 ATOM						61.866	-3.697	70.678	1.00 26.54
ATOM 5242 CA ARN B 280 64,318 -3.046 69,680 1.00 23.70 ATOM 5244 CG ARN B 280 65,520 -2.248 69,164 1.00 22.63 ATOM 5245 OD1 ARN B 280 65,107 -0.937 68,505 1.00 30.83 ATOM 5246 ND2 ARN B 280 65,107 -0.937 68,505 1.00 30.83 ATOM 5246 ND2 ARN B 280 65,107 -0.937 68,505 1.00 30.83 ATOM 5246 ND2 ARN B 280 65,107 -0.937 68,505 1.00 25,63 ATOM 5247 C ARN B 280 64,775 -5.321 69,124 1.00 26,16 ATOM 5248 O ARN B 280 64,775 -5.321 69,124 1.00 26,16 ATOM 5248 O ARN B 280 64,775 -5.321 69,124 1.00 26,16 ATOM 5250 CA ILE B 281 65,485 -6.067 71,667 1.00 25,18 ATOM 5250 CA ILE B 281 66,046 -7.527 71,667 1.00 25,18 ATOM 5251 CB ILE B 281 66,046 -7.527 73,648 1.00 28,53 ATOM 5252 CG2 ILE B 281 66,046 -7.527 73,648 1.00 28,53 ATOM 5253 CG1 ILE B 281 66,046 -7.527 73,648 1.00 28,53 ATOM 5255 C ILE B 281 66,046 -7.527 73,648 1.00 28,53 ATOM 5255 C ILE B 281 66,036 -7.527 73,648 1.00 28,53 ATOM 5255 C ILE B 281 66,036 -7.527 73,648 1.00 28,53 ATOM 5255 C ILE B 281 66,036 -7.527 73,648 1.00 28,53 ATOM 5255 C ILE B 281 64,320 -7.030 71,507 1.00 25,77 ATOM 5255 C ILE B 281 64,320 -7.030 71,507 1.00 25,77 ATOM 5255 C VAL B 282 63,139 -6,618 71,950 1.00 21,30 ATOM 5257 N VAL B 282 63,139 -6,618 71,950 1.00 21,30 ATOM 5258 CA VAL B 282 63,1691 -7,465 71,813 1.00 22,07 ATOM 5260 CG1 VAL B 282 60,763 -6,775 72,387 1.00 24,07 ATOM 5261 CG2 VAL B 282 60,763 -6,775 72,387 1.00 24,07 ATOM 5262 C VAL B 282 61,789 -7,795 70,339 1.00 24,07 ATOM 5262 C VAL B 282 61,462 -7,795 70,339 1.00 24,07 ATOM 5262 C VAL B 282 61,462 -7,795 70,339 1.00 24,07 ATOM 5264 N ARG B 283 61,570 -6,779 69,488 1.00 23,87 ATOM 5266 CB ARG B 283 61,500 -5,677 73,906 1.00 22,87 ATOM 5267 C ARG B 283 60,210 -5,612 67,399 1.00 25,48 ATOM 5268 CD ARG B 283 60,379 -7,795 69,488 1.00 23,87 ATOM 5268 CD ARG B 283 60,379 -7,795 69,488 1.00 23,87 ATOM 5268 CD ARG B 283 60,379 -7,795 69,488 1.00 23,87 ATOM 5270 CZ ARG B 283 60,379 -7,795 69,488 1.00 23,87 ATOM 5270 CZ ARG B 283 60,379 -7,795 69,488 1.00 23,87 ATOM 5280 PC VAL B 285 60,685 -7,796 69,385 1.00 29,76 ATOM 5290 CD PLE B								70.804	1.00 21.93
ATOM 5243 CB ASN B 280 65.520 -2.248 69.164 1.00 22.63 ATOM 5245 ODI ASN B 280 65.900 0.112 68.7796 1.00 23.81 ATOM 5246 ND2 ASN B 280 65.900 0.112 68.714 1.00 22.63 ATOM 5246 ND2 ASN B 280 64.746 -4.466 70.009 1.00 26.10 ATOM 5248 0 ASN B 280 64.7746 -4.466 70.009 1.00 26.10 ATOM 5248 0 ASN B 280 64.7746 -5.321 69.124 1.00 26.15 ATOM 5249 N ILE B 281 65.080 -4.724 71.272 1.00 26.15 ATOM 5259 N ILE B 281 66.006 -6.009 71.667 1.00 25.81 ATOM 5251 CB ILE B 281 66.006 -6.009 73.124 1.00 28.53 ATOM 5251 CB ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5252 CG2 ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 CC ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 CC ILE B 281 66.006 7.527 73.648 1.00 28.24 ATOM 5255 CC ILE B 281 66.006 7.527 73.648 1.00 28.24 ATOM 5255 CC ILE B 281 64.484 -8.131 70.992 1.00 23.39 ATOM 5255 C ILE B 281 64.484 -8.131 70.992 1.00 23.39 ATOM 5255 C ILE B 281 64.484 -8.131 70.992 1.00 23.39 ATOM 5255 C ILE B 281 64.484 -8.131 70.992 1.00 23.39 ATOM 5258 CA VAL B 282 63.139 -6.618 71.950 1.00 22.70 ATOM 5258 CA VAL B 282 60.703 -7.657 72.387 1.00 24.90 ATOM 5259 CC VAL B 282 60.703 -7.677 72.387 1.00 24.90 ATOM 5250 CG1 VAL B 282 60.703 -7.677 72.387 1.00 22.90 ATOM 5250 CG1 VAL B 282 60.656 -6.587 73.906 1.00 22.38 ATOM 5260 CG1 VAL B 282 61.662 -6.578 73.906 1.00 22.38 ATOM 5265 CA ARG 8 283 61.576 -6.971 68.060 1.00 26.85 ATOM 5265 CA ARG 8 283 61.576 -6.971 68.060 1.00 26.85 ATOM 5265 CA ARG 8 283 61.576 -6.971 68.060 1.00 26.85 ATOM 5265 CA ARG 8 283 61.576 -6.971 68.060 1.00 26.45 ATOM 5266 CB ARG 8 283 61.576 -6.971 68.060 1.00 26.38 ATOM 5267 CA ARG 8 283 61.576 -6.971 68.060 1.00 26.48 ATOM 5267 CA ARG 8 283 61.576 -6.971 68.060 1.00 26.48 ATOM 5267 CA ARG 8 283 60.370 -7.606 67.992 1.00 25.48 ATOM 5268 CA ARG 8 283 60.370 -7.606 67.992 1.00 25.48 ATOM 5267 CA ARG 8 283 60.370 -7.821 69.993 1.00 25.48 ATOM 5270 CZ ARG 8 283 60.370 -7.821 69.994 1.00 26.43 ATOM 5270 CZ ARG 8 283 60.370 -7.821 69.994 1.00 26.43 ATOM 5270 CZ ARG 8 283 60.370 -7.821 69.994 1.00 26.43 ATOM 5270 CZ									
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ATOM 5246 ND2 ASN B 280 64.094 -0.878 67.796 1.00 25.84 ATOM 5247 C ASN B 280 65.900 0.112 68.714 1.00 26.54 ATOM 5248 O ASN B 280 64.746 -4.466 70.009 1.00 26.10 ATOM 5249 N ILE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5249 N ILE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5250 CA ILE B 281 66.006 -6.098 73.124 1.00 26.51 ATOM 5251 CB ILE B 281 66.006 -7.527 73.648 1.00 28.50 ATOM 5252 CC2 ILE B 281 66.006 -7.527 73.648 1.00 28.50 ATOM 5253 CG1 ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 CC1 ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 66.006 -7.527 73.648 1.00 28.24 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 23.77 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 23.77 ATOM 5256 O ILE B 281 64.320 -7.030 71.507 1.00 23.77 ATOM 5256 C ILE B 282 661.961 -7.465 71.813 1.00 24.90 ATOM 5257 N VAL B 282 63.139 -6.618 71.950 1.00 21.39 ATOM 5258 CA VAL B 282 60.703 -6.775 72.387 1.00 24.02 ATOM 5250 CG1 VAL B 282 60.865 6.587 73.906 1.00 22.38 ATOM 5261 CG2 VAL B 282 60.865 6.687 73.906 1.00 22.88 ATOM 5262 C VAL B 282 61.718 -7.795 70.339 1.00 22.28 ATOM 5265 C VAL B 282 61.718 -7.795 70.339 1.00 22.28 ATOM 5265 C ARG B 283 61.576 -5.971 68.060 1.00 26.83 ATOM 5265 C ARG B 283 61.576 -6.971 68.060 1.00 26.83 ATOM 5265 C ARG B 283 61.576 -6.971 68.060 1.00 26.85 ATOM 5267 C G ARG B 283 60.337 -4.760 67.838 1.00 24.43 ATOM 5270 NE ARG B 283 60.337 -4.760 67.838 1.00 24.43 ATOM 5271 NH1 ARG B 283 60.634 -7.855 67.402 1.00 25.48 ATOM 5272 NH2 ARG B 283 60.634 -7.855 67.402 1.00 25.48 ATOM 5270 C C ARG B 283 60.634 -7.855 67.402 1.00 25.48 ATOM 5271 NH1 ARG B 283 60.634 -7.855 67.402 1.00 25.48 ATOM 5272 NH2 ARG B 283 60.634 -7.855 67.402 1.00 25.48 ATOM 5273 C C ARG B 283 60.634 -7.855 67.402 1.00 25.48 ATOM 5276 C C ARG B 283 60.634 -7.855 67.402 1.00 28.43 ATOM 5277 NH1 ARG B 283 60.634 -7.855 67.402 1.00 28.43 ATOM 5278 C C C C C C C C C C C C C C C C C C C	MOTA	5244	CG	ASN B	280	65.107	-0.937		
ATOM 5246 ND2 ASN B 280 65.900 0.112 68.714 1.00 26.54 ATOM 5247 C ASN B 280 64.746 -4.466 70.009 1.00 26.16 ATOM 5248 0 ASN B 280 64.775 -5.321 69.124 1.00 26.16 ATOM 5248 0 ASN B 280 65.980 -4.724 71.272 1.00 26.16 ATOM 5250 CQ ILLE B 281 65.485 -6.067 71.667 1.00 26.16 ATOM 5251 CB ILLE B 281 66.046 -7.527 73.648 1.00 28.53 ATOM 5252 CQ2 ILLE B 281 66.046 -7.527 73.648 1.00 28.53 ATOM 5253 CG1 ILLE B 281 66.046 -7.527 73.648 1.00 28.53 ATOM 5255 CI ILLE B 281 66.036 -5.442 74.571 1.00 22.24 ATOM 5254 CD1 ILLE B 281 66.036 -7.407 71.507 1.00 22.74 ATOM 5255 C ILLE B 281 64.320 -7.030 71.507 1.00 23.73 ATOM 5255 C ILLE B 281 64.320 -7.030 71.507 1.00 23.73 ATOM 5255 C ILLE B 281 64.320 -7.030 71.507 1.00 23.73 ATOM 5256 0 ILLE B 282 61.961 -7.465 71.813 1.00 22.74 ATOM 5256 0 ILLE B 282 61.961 -7.465 71.813 1.00 22.74 ATOM 5258 C AVAL B 282 61.961 -7.465 71.813 1.00 22.74 ATOM 5258 C AVAL B 282 61.961 -7.465 71.813 1.00 22.07 ATOM 5259 CB VAL B 282 61.961 -7.465 71.813 1.00 22.07 ATOM 5250 CC1 VAL B 282 60.703 -6.775 72.387 1.00 24.07 ATOM 5261 CC2 VAL B 282 61.961 -7.465 71.813 1.00 22.65 ATOM 5263 C VAL B 282 61.961 -7.465 71.813 1.00 22.65 ATOM 5265 C ARG B 283 61.500 -7.79 69.948 1.00 23.75 ATOM 5265 C ARG B 283 61.500 -7.95 99.78 1.00 22.65 ATOM 5266 CB ARG B 283 61.500 -7.95 69.978 1.00 22.65 ATOM 5266 CB ARG B 283 61.500 -7.600 67.838 1.00 22.65 ATOM 5266 CB ARG B 283 60.337 -4.760 67.838 1.00 22.65 ATOM 5267 CG ARG B 283 60.337 -4.760 67.838 1.00 22.65 ATOM 5268 CD ARG B 283 60.337 -4.760 67.838 1.00 22.65 ATOM 5268 CD ARG B 283 60.337 -4.760 67.838 1.00 22.65 ATOM 5267 CG ARG B 283 60.337 -4.760 67.838 1.00 22.65 ATOM 5268 CD ARG B 283 60.340 -2.356 63.825 1.00 22.64 ATOM 5270 CZ ARG B 283 60.360 -7.4760 67.838 1.00 22.65 ATOM 5270 CZ ARG B 283 60.360 -7.4760 67.838 1.00 22.65 ATOM 5270 CZ ARG B 283 60.360 -7.4760 67.838 1.00 22.65 ATOM 5270 CZ ARG B 283 60.360 -7.4760 67.838 1.00 22.65 ATOM 5270 CZ ARG B 283 60.360 -7.500 -7.600 67.33 67.339 1.00 23.75 ATOM 5270 CZ ARG B 283 60.360 -7.500 -7.600 67.3			OD1	ASN B	280	64.094	-0.878	67.796	1.00 25.81
ATOM 5247 C ASN B 280 64.746 -4.466 70.009 1.00 26.10 ATOM 5248 N SUB 280 64.775 -5.321 69.124 1.00 26.10 ATOM 5259 N ILE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5250 CA ILE B 281 66.086 -6.098 73.124 1.00 28.50 ATOM 5251 CE ILE B 281 66.006 -7.527 73.648 1.00 28.50 ATOM 5252 CC2 ILE B 281 66.006 -7.527 73.648 1.00 28.50 ATOM 5253 CC1 ILE B 281 66.006 -7.527 73.648 1.00 28.50 ATOM 5253 CC1 ILE B 281 66.006 -7.527 73.648 1.00 28.50 ATOM 5255 C ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 25.74 ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 25.74 ATOM 5256 O ILE B 281 64.484 -8.131 70.982 1.00 23.39 ATOM 5257 N VAL B 282 63.139 -6.618 71.950 1.00 21.39 ATOM 5258 CA VAL B 282 66.703 -7.465 71.813 1.00 22.90 ATOM 5250 CG1 VAL B 282 60.703 -6.775 72.387 1.00 24.02 ATOM 5260 CG1 VAL B 282 60.865 -6.587 73.906 1.00 22.38 ATOM 5261 CG2 VAL B 282 60.865 -6.587 73.906 1.00 22.38 ATOM 5262 C VAL B 282 61.718 -7.795 70.339 1.00 22.28 ATOM 5263 O VAL B 282 61.718 -7.795 70.339 1.00 22.58 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 26.387 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 26.55 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 26.55 ATOM 5267 CG ARG B 283 60.337 -4.760 67.838 1.00 22.19 ATOM 5268 CD ARG B 283 60.374 -4.760 67.838 1.00 22.15 ATOM 5267 CG ARG B 283 60.375 -4.760 65.622 1.00 29.64 ATOM 5270 NE ARG B 283 60.442 -3.333 67.339 1.00 31.50 ATOM 5268 CD ARG B 283 60.442 -3.350 63.805 1.00 26.55 ATOM 5267 CG ARG B 283 60.444 -7.815 69.908 1.00 26.48 ATOM 5270 NE ARG B 283 60.444 -7.855 67.902 1.00 26.48 ATOM 5271 NH1 ARG B 283 60.634 -7.855 67.402 1.00 26.44 ATOM 5273 C ARG B 283 60.634 -7.855 67.402 1.00 26.44 ATOM 5275 N QUB 284 66.289 -8.260 67.992 1.00 38.19 ATOM 5268 CD ARG B 283 60.634 -7.855 67.402 1.00 28.94 ATOM 5270 C ARG B 283 60.634 -7.855 67.402 1.00 28.94 ATOM 5270 C ARG B 283 60.634 -7.855 67.402 1.00 28.94 ATOM 5280 C C ARG B 283						65.900	0.112	68.714	1.00 26.54
ATOM 5248 O ASN B 280							-4.466	70.009	1.00 26.10
ATOM 5259 CA ILE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5250 CA ILE B 281 65.080 -4.724 71.272 1.00 26.10 ATOM 5251 CB ILE B 281 66.006 -6.098 73.124 1.00 28.51 ATOM 5252 CG2 ILE B 281 66.006 -7.527 73.648 1.00 28.53 ATOM 5253 CG1 ILE B 281 67.392 -5.447 73.671 1.00 32.07 ATOM 5254 CD1 ILE B 281 68.038 -5.442 74.541 1.00 28.24 ATOM 5255 C ILE B 281 64.484 -8.131 70.982 1.00 23.39 ATOM 5255 C ILE B 281 64.484 -8.131 70.982 1.00 23.39 ATOM 5257 N VAL B 282 63.139 -6.618 71.995 1.00 23.39 ATOM 5258 CA VAL B 282 60.703 -6.775 72.387 1.00 24.07 ATOM 5259 CB VAL B 282 60.703 -6.775 72.387 1.00 24.07 ATOM 5250 CG1 VAL B 282 59.464 -7.611 72.093 1.00 22.28 ATOM 5261 CG2 VAL B 282 60.865 -6.587 73.906 1.00 26.89 ATOM 5261 CG2 VAL B 282 61.462 -8.949 69.978 1.00 23.87 ATOM 5262 C VAL B 282 61.462 -8.949 69.978 1.00 23.87 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 22.58 ATOM 5265 CB ARG B 283 61.576 -6.971 68.060 1.00 22.54 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 22.54 ATOM 5265 CB ARG B 283 61.576 -6.971 68.060 1.00 22.54 ATOM 5265 CB ARG B 283 61.576 -6.971 68.060 1.00 22.54 ATOM 5267 CB ARG B 283 60.337 -4.760 67.838 1.00 22.54 ATOM 5268 CD ARG B 283 60.337 -4.760 67.838 1.00 22.54 ATOM 5267 CB ARG B 283 60.337 -4.760 67.838 1.00 22.54 ATOM 5268 CD ARG B 283 60.337 -4.760 67.838 1.00 22.54 ATOM 5267 CB ARG B 283 60.337 -4.760 67.838 1.00 22.54 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5268 CD ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5268 CD ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5270 CZ ARG B 283 60.442 -3.333 60.442 -3.356 66.382 1.00 32.442 ATOM 5270 CZ ARG B 283 60.464 CZ ARG B 283 60.464									
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ATOM 5255 C ILE B 281 68.038 -5.442 74.541 1.00 28.27 ATOM 5256 C ILE B 281 64.320 -7.030 71.507 1.00 25.77 ATOM 5257 N VAL B 282 63.139 -6.618 71.950 1.00 23.39 ATOM 5258 CA VAL B 282 61.961 -7.465 71.813 1.00 22.90 ATOM 5259 CB VAL B 282 60.703 -6.775 72.387 1.00 24.93 ATOM 5260 CG1 VAL B 282 60.703 -6.775 72.387 1.00 22.28 ATOM 5261 CG2 VAL B 282 60.703 -6.775 72.387 1.00 22.28 ATOM 5262 C VAL B 282 61.961 -7.465 71.813 1.00 22.28 ATOM 5263 O VAL B 282 61.766 -6.587 73.906 1.00 22.88 ATOM 5264 N ARG B 283 61.799 -6.799 69.488 1.00 23.87 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 27.95 ATOM 5266 CB ARG B 283 61.576 -6.971 68.060 1.00 27.95 ATOM 5267 CG ARG B 283 60.337 -4.760 67.838 1.00 26.55 ATOM 5268 CD ARG B 283 60.337 -4.760 67.838 1.00 26.55 ATOM 5269 NE ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5270 CZ ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5271 NH1 ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5272 NH2 ARG B 283 62.341 -8.552 66.31 1.00 29.64 ATOM 5273 C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5274 O ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5276 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 31.50 ATOM 5278 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5278 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5276 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5276 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5276 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5278 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5278 C C ARG B 283 62.341 -8.552 66.31 1.00 29.76 ATOM 5279 C D GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5280 C C BLU B 284 66.299 -8.260 67.992 1.00 38.31 ATOM 5281 C C C ARG B 283 63.667 -11.802 7.0663 1.00 29.84 ATOM 5280 C C BLU B 284 66.099 1.00 4.899 1.00 25.71 ATOM 5280 C C BLU B 284 66.090 -1.1400 66.913 1.00 28.09 ATOM 5280 C C BLU B 284 66.090 -1.1400 66.913 1.00 28.09 ATOM 5280 C C BLU B 285 62.460 -12.25 68.758 1.00 31.93 ATOM 5280 C C BLU B 285 62.460 -12.25 68.758 1.00 31.93 ATOM 528			CG1	ILE B	281	67.392	-5.454	73. 17 3	1.00 32.07
ATOM 5255 C ILE B 281 64.320 -7.030 71.507 1.00 25.777 ATOM 5256 O ILE B 281 64.484 -8.131 70.982 1.00 23.39 ATOM 5257 N VAL B 282 61.961 -7.465 71.813 1.00 22.98 ATOM 5258 CA VAL B 282 61.961 -7.465 71.813 1.00 24.07 ATOM 5250 CG VAL B 282 60.703 -6.618 71.950 1.00 24.07 ATOM 5260 CG1 VAL B 282 60.865 -6.587 73.996 1.00 24.07 ATOM 5261 CG2 VAL B 282 60.865 -6.587 73.996 1.00 24.87 ATOM 5262 C VAL B 282 61.788 -7.795 70.339 1.00 23.87 ATOM 5263 O VAL B 282 61.789 -7.795 70.339 1.00 23.87 ATOM 5264 N ARG B 283 61.576 -6.971 68.060 1.00 23.19 ATOM 5265 CA ARG B 283 61.579 -6.779 69.488 1.00 23.19 ATOM 5266 CB ARG B 283 61.510 -5.612 67.359 1.00 25.48 ATOM 5266 CB ARG B 283 60.337 -4.760 67.838 1.00 25.55 ATOM 5266 CB ARG B 283 60.337 -4.760 67.838 1.00 25.55 ATOM 5268 N C ARG B 283 60.337 -4.760 67.838 1.00 25.55 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 25.55 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 25.55 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 25.55 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 25.55 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 25.55 ATOM 5270 CZ ARG B 283 60.915 -2.414 65.116 1.00 26.604 ATOM 5271 NH1 ARG B 283 60.634 -2.356 63.825 1.00 29.64 ATOM 5273 C ARG B 283 62.604 -7.855 67.402 1.00 32.04 ATOM 5276 CA GLU B 284 66.934 -8.552 66.431 1.00 29.76 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 38.93 ATOM 5278 C GLU B 284 66.289 -8.260 67.992 1.00 38.93 ATOM 5280 CE GLU B 284 66.899 -10.300 68.901 1.00 28.09 ATOM 5281 CC GLU B 284 66.934 -8.566 67.381 1.00 29.64 ATOM 5282 C GLU B 284 64.699 -10.300 68.901 1.00 28.09 ATOM 5280 CE GLU B 284 64.699 -10.300 68.901 1.00 28.09 ATOM 5281 CC GLU B 284 64.699 -10.300 68.901 1.00 28.09 ATOM 5282 C GLU B 284 64.699 -10.300 68.901 1.00 28.09 ATOM 5280 CE JUB 284 66.798 -6.548 67.901 1.00 28.09 ATOM 5280 CE JUB 284 66.798 -6.666 67.381 1.00 31.19 ATOM 5280 C GLU B 284 66.798 -6.666 67.381 1.00 32.42 ATOM 5290 C VAL B 285 63.687 -11.802 70.863 1.00 28.93 ATOM 5290 C VAL B 285 63.687 -11.802 70.863 1.00 28.93 ATOM 5290 C PHE						68.038	-5.442	74.541	1.00 28.24
ATOM 5256								71.507	1.00 25.77
ATOM 5257 N VAL B 282 63 139 -6.618 71.950 1.00 21.30 ATOM 5258 CA VAL B 282 61.961 -7.465 71.813 1.00 22.90 ATOM 5259 CB VAL B 282 60.703 -6.775 72.387 1.00 24.07 ATOM 5260 CG1 VAL B 282 60.703 -6.775 72.387 1.00 24.07 ATOM 5261 CG2 VAL B 282 60.865 -6.587 73.906 1.00 22.87 ATOM 5263 O VAL B 282 61.462 -8.949 69.978 1.00 23.87 ATOM 5263 O VAL B 282 61.462 -8.949 69.978 1.00 23.87 ATOM 5264 N ARG B 283 61.576 -6.971 68.060 1.00 27.85 ATOM 5266 CB ARG B 283 61.576 -6.971 68.060 1.00 27.58 ATOM 5266 CB ARG B 283 61.510 -5.612 67.359 1.00 25.48 ATOM 5266 CB ARG B 283 60.337 -4.760 67.838 1.00 26.55 ATOM 5269 NE ARG B 283 60.210 -3.208 65.908 1.00 24.43 ATOM 5269 NE ARG B 283 60.210 -3.208 65.908 1.00 24.43 ATOM 5270 CZ ARG B 283 60.915 -2.444 65.116 1.00 26.04 ATOM 5271 NH1 ARG B 283 60.915 -2.444 65.116 1.00 26.04 ATOM 5273 C ARG B 283 60.634 -2.356 63.825 1.00 29.04 ATOM 5273 C ARG B 283 62.341 -8.552 66.431 1.00 29.76 ATOM 5276 CA GLU B 284 63.859 -7.821 67.923 1.00 31.04 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 32.04 ATOM 5276 CA GLU B 284 66.289 -8.260 67.992 1.00 32.04 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 32.04 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 32.04 ATOM 5278 CG GLU B 284 66.289 -8.260 67.992 1.00 38.93 ATOM 5280 CE1 GLU B 284 66.289 -8.260 67.992 1.00 38.93 ATOM 5280 CE1 GLU B 284 66.798 -6.864 67.601 1.00 26.54 ATOM 5280 CE GLU B 284 66.099 -10.340 68.901 1.00 32.42 ATOM 5281 CC2 GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CE1 GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CC GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CC GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CC GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CC GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CC GLU B 284 66.099 -10.340 68.901 1.00 32.80 ATOM 5280 CC GLU B 285 65.037 -11.470 71.478 1.00 32.82 ATOM 5280 CC VAL B 285 63.687 -11.802 70.863 1.00 25.95 ATOM 5280 CC VAL B 285 65.037 -11.400 71.479 1.00 32.83 ATOM 5290 C VAL B 285 65.906 -11.800 70.91 1.00 25.51				TIE D	201				
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ATOM 5261 CG2 VAL B 282 60.865 -6.587 73.906 1.00 26.89 ATOM 5263 0 VAL B 282 61.718 -7.795 70.339 1.00 23.87 ATOM 5263 0 VAL B 282 61.718 -7.795 70.339 1.00 23.87 ATOM 5265 CA ARG B 283 61.579 -6.779 69.488 1.00 23.19 ATOM 5265 CA ARG B 283 61.570 -6.971 68.060 1.00 27.95 ATOM 5266 CB ARG B 283 61.510 -5.612 67.359 1.00 25.48 ATOM 5266 CB ARG B 283 60.337 -4.760 67.838 1.00 26.55 ATOM 5268 CD ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5269 NE ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5271 NH1 ARG B 283 60.915 -2.414 65.116 1.00 26.44 ATOM 5271 NH1 ARG B 283 60.915 -2.414 65.116 1.00 26.44 ATOM 5271 NH1 ARG B 283 60.915 -2.414 65.116 1.00 26.44 ATOM 5271 NH2 ARG B 283 60.634 -2.356 63.825 1.00 29.64 ATOM 5273 C ARG B 283 62.341 -8.552 66.431 1.00 29.76 ATOM 5275 N GLU B 284 63.859 -7.821 67.923 1.00 31.50 ATOM 5275 N GLU B 284 63.859 -7.821 67.923 1.00 31.50 ATOM 5277 CB GLU B 284 66.798 -6.864 67.381 1.00 29.76 ATOM 5277 CB GLU B 284 66.798 -6.864 67.992 1.00 38.31 ATOM 5280 OEI GLU B 284 66.798 -6.864 67.992 1.00 38.31 ATOM 5280 OEI GLU B 284 66.798 -6.864 67.640 1.00 48.93 ATOM 5280 OEI GLU B 284 66.798 -6.864 67.640 1.00 48.93 ATOM 5280 OEI GLU B 284 66.836 -10.105 67.714 1.00 31.93 ATOM 5280 OEI GLU B 284 66.838 -10.105 67.714 1.00 31.93 ATOM 5280 OEI GLU B 284 66.838 -10.105 67.714 1.00 31.93 ATOM 5280 OEI GLU B 284 66.899 -11.001 66.913 1.00 28.26 ATOM 5285 CA VAL B 285 63.687 -11.802 70.863 1.00 28.26 ATOM 5280 CB UAL B 285 63.687 -11.802 70.863 1.00 28.25 ATOM 5280 CB UAL B 285 63.687 -11.802 70.863 1.00 28.25 ATOM 5292 CA WAL B 285 63.687 -11.802 70.863 1.00 28.25 ATOM 5292 CA WAL B 285 63.687 -11.802 70.863 1.00 28.25 ATOM 5292 CA WAL B 285 63.687 -11.802 70.863 1.00 28.25 ATOM 5292 CA WAL B 285 63.687 -11.802 70.863 1.00 28.25 ATOM 5293 CB PHE B 286 60.105 -11.948 68.299 1.00 28.25 ATOM 5292 CA WAL B 285 63.687 -11.800 70.863 1.00 28.25 ATOM 5299 C WAL B 285 63.687 -11.800 70.863 1.00 28.25 ATOM 5299 CP HE B 286 69.969 -10.00 55.16 ATOM		5260	CG1	VAL B	282	59.464	-7.611		
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ATOM 5265 CA ARG B 283 61.799 -6.779 69.488 1.00 23.19 ATOM 5265 CA ARG B 283 61.576 -6.971 68.060 1.00 27.95 ATOM 5266 CB ARG B 283 61.576 -6.971 68.060 1.00 27.95 ATOM 5266 CB ARG B 283 60.337 -4.760 67.838 1.00 25.48 ATOM 5269 NE ARG B 283 60.442 -3.333 67.339 1.00 31.52 ATOM 5269 NE ARG B 283 60.210 -3.208 65.908 1.00 24.43 ATOM 5270 CZ ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5271 NH1 ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5272 NH2 ARG B 283 60.915 -2.414 65.116 1.00 26.04 ATOM 5273 C ARG B 283 60.634 -2.356 63.825 1.00 29.64 ATOM 5273 C ARG B 283 62.634 -7.855 67.402 1.00 32.04 ATOM 5275 N GLU B 284 63.859 -7.821 67.923 1.00 31.50 ATOM 5276 CA GLU B 284 66.289 -8.260 67.992 1.00 32.42 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5278 CG GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5279 CD GLÜ B 284 68.102 -6.518 68.362 1.00 57.37 ATOM 5280 OE1 GLU B 284 68.102 -6.518 68.362 1.00 57.37 ATOM 5281 OE2 GLU B 284 68.102 -6.518 68.362 1.00 57.37 ATOM 5283 O GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5284 N VAL B 285 64.089 -10.340 68.901 1.00 28.26 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.26 ATOM 5287 CGI VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5288 CG VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.39 ATOM 5289 C VAL B 285 63.687 -11.802 70.863 1.00 28.21 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.39 ATOM 5291 CD PHE B 286 69.064 -11.853 69.374 1.00 24.57 ATOM 5292 CA PHE B 286 69.064 -11.853 69.374 1.00 24.57 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5299 CZ PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5299 CD PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5299 CD PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5299 CD PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5299 CD PHE B 286 59.072 -14.180 70.319 1.00 25.16 ATOM 5299 CD PHE B 286 59.072 -14.180 70.319 1.00 25.16 ATOM 5299 CD PHE B 286 59.072 -14.578 72.626 1.00 25								69.978	1.00 22.65
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ATOM 5270 CZ ARG B 283 60.915 -2.414 65.116 1.00 26.45 ATOM 5271 NH1 ARG B 283 61.902 -1.676 65.622 1.00 26.04 ATOM 5272 NH2 ARG B 283 60.634 -2.356 63.825 1.00 29.64 ATOM 5273 C ARG B 283 62.634 -7.855 67.402 1.00 32.04 ATOM 5274 O ARG B 283 62.341 -8.552 66.431 1.00 29.76 ATOM 5275 N GLU B 284 63.859 -7.821 67.923 1.00 31.50 ATOM 5276 CA GLU B 284 64.934 -8.646 67.381 1.00 32.42 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5278 CG GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5279 CD GLU B 284 68.102 -6.518 68.362 1.00 55.28 ATOM 5280 OE1 GLU B 284 68.102 -6.518 68.362 1.00 55.28 ATOM 5280 OE1 GLU B 284 68.102 -6.518 68.362 1.00 55.42 ATOM 5280 OE1 GLU B 284 66.889 -11.00 66.913 1.00 28.26 ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5285 CB VAL B 285 63.657 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5287 CGI VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5289 C VAL B 285 63.687 -11.470 71.478 1.00 26.93 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.38 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.39 ATOM 5291 N PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5296 CD PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.064 -11.853 69.374 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5299 CC PHE B 286 59.664 -11.853 69.374 1.00 22.51 ATOM 5299 CC PHE B 286 59.664 -11.853 69.374 1.00 22.51 ATOM 5299 CC PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5299 CC PHE B 286 59.651 -12.331 72.833 1.00 22.92 ATOM 5299 CC PHE B 286 59.651 -12.331 72.833 1.00 22.92 ATOM 5299 CC PHE B 286 59.651 -12.331 72.833 1.00 22.59 ATOM 5299 CC PHE B 286 59.651 -12.331 72.833 1.00 22.59 ATOM 5299 CC PHE B 286 59.651 -12.331 72.833 1.00 22.59 ATOM 5299 CC PHE B 286 59.651 -12.331 72.833 1.00 22.59 ATOM 5300 C PHE B 286 59.5772 -14.578 72.666 1.00 22.51 ATOM 5300 C PHE B 286 59.588 -11.630 66.620		5269	NE	ARG B	283	60.210	-3.208	65.908	
ATOM 5271 NH1 ARG B 283 61.902 -1.676 65.622 1.00 26.04 ATOM 5272 NH2 ARG B 283 60.634 -2.356 63.825 1.00 29.64 ATOM 5273 C ARG B 283 62.634 -7.855 67.402 1.00 32.04 ATOM 5274 O ARG B 283 62.541 -8.552 66.431 1.00 29.76 ATOM 5275 N GLU B 284 63.859 -7.821 67.923 1.00 31.50 ATOM 5276 CA GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5277 CB GLU B 284 66.289 -8.260 67.992 1.00 38.31 ATOM 5279 CD GLU B 284 66.798 -6.864 67.640 1.00 48.93 ATOM 5279 CD GLU B 284 68.102 -6.518 68.362 1.00 56.28 ATOM 5280 OE1 GLU B 284 68.102 -6.518 68.362 1.00 55.42 ATOM 5281 OE2 GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.69.94 -7.281 68.901 1.00 28.26 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.26 ATOM 5287 CGI VAL B 285 63.667 -11.400 71.478 1.00 28.33 ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.38 ATOM 5291 N PHE B 286 60.105 -11.948 68.249 1.00 24.57 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 60.05 -14.180 69.374 1.00 22.92 ATOM 5293 CB PHE B 286 69.064 -11.853 69.374 1.00 22.51 ATOM 5296 CD2 PHE B 286 69.05 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 286 69.05 -14.180 70.319 1.00 22.51 ATOM 5298 CC2 PHE B 286 69.05 -14.180 70.319 1.00 22.51 ATOM 5299 CC PHE B 286 69.05 -14.180 70.319 1.00 22.51 ATOM 5299 CC PHE B 286 69.05 -14.180 70.319 1.00 22.51 ATOM 5299 CC PHE B 286 69.05 -14.180 70.319 1.00 22.51 ATOM 5290 CC PHE B 286 69.59.518 -11.318 66.993 1.00 22.590 ATOM 5290 CC PHE B 286 69.59.518 -11.318 66.993 1.00 22.590 ATOM 5300 C PHE B 286 69.59.518 -11.318 66.993 1.00 22.849 ATOM 5300 C PHE B 286 69.59.518 -11.318 66.993 1.00 22.590 ATOM 5301 O PHE B 286 69.772 -14.578 72.666 1.00 26.75 ATOM 5300 C PHE B 286 69.772 -14.578 72.666 1.00 22.849 ATOM 5300 C PHE B 286 69.59.518 -11.318 66.329 1.00 22.829				ARG B	283	60.915	-2.414	65.116	1.00 26.45
ATOM 5272 NH2 ARG B 283 60.634 -2.356 63.825 1.00 29.64 ATOM 5273 C ARG B 283 62.634 -7.855 67.402 1.00 32.04 ATOM 5274 O ARG B 283 62.341 -8.552 66.431 1.00 29.76 ATOM 5275 N GLU B 284 63.859 -7.821 67.923 1.00 31.50 ATOM 5276 CA GLU B 284 66.298 -8.260 67.381 1.00 32.42 ATOM 5277 CB GLU B 284 66.299 -8.260 67.992 1.00 38.31 ATOM 5278 CG GLU B 284 66.798 -6.864 67.640 1.00 48.93 ATOM 5279 CD GLU B 284 66.798 -6.864 67.640 1.00 48.93 ATOM 5279 CD GLU B 284 68.102 -6.518 68.362 1.00 56.28 ATOM 5280 OE1 GLU B 284 68.102 -6.518 68.222 1.00 55.42 ATOM 5281 OE2 GLU B 284 68.150 -5.485 69.069 1.00 55.42 ATOM 5282 C GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.257 -13.206 71.262 1.00 29.84 ATOM 5289 C VAL B 285 65.037 -11.470 71.478 1.00 28.33 ATOM 5280 CG2 VAL B 285 65.037 -11.470 71.478 1.00 28.33 ATOM 5290 O VAL B 285 62.420 -12.265 68.758 1.00 31.19 ATOM 5290 C PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5291 N PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5292 CA PHE B 286 60.105 -11.948 69.374 1.00 22.51 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.57 ATOM 5295 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.57 ATOM 5295 CD2 PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5298 CE2 PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.551 -12.331 71.779 1.00 25.51 ATOM 5299 CZ PHE B 286 59.518 -11.318 66.993 1.00 22.92 ATOM 5300 C PHE B 286 59.518 -11.578 66.993 1.00 22.92 ATOM 5300 C PHE B 286 59.518 -11.578 66.993						61.902	-1.676	65.622	1.00 26.04
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ATOM 5280 OE1 GLU B 284 69.084 -7.281 68.222 1.00 57.37 ATOM 5281 OE2 GLU B 284 68.150 -5.485 69.069 1.00 55.42 ATOM 5282 C GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5285 CA VAL B 285 64.089 -10.340 68.901 1.00 28.09 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5287 CG1 VAL B 285 63.257 -13.206 71.262 1.00 29.84 ATOM 5288 CG2 VAL B 285 65.037 -11.470 71.478 1.00 26.93 ATOM 5289 C VAL B 285 65.037 -11.470 71.478 1.00 26.93 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD1 PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.92 ATOM 5298 CE2 PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5298 CE2 PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5298 CE2 PHE B 286 59.772 -14.578 72.626 1.00 22.92 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 22.92 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 22.84 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 22.84 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.338 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.338 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 22.84 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 23.338 ATOM 5300 C PHE B 286 59.772 -1			CD	GLŬ B	284	68.102	-6.518	68.362	
ATOM 5281 OE2 GLU B 284 68.150 -5.485 69.069 1.00 55.42 ATOM 5282 C GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5284 N VAL B 285 64.089 -10.340 68.901 1.00 28.09 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5287 CG1 VAL B 285 63.257 -13.206 71.262 1.00 29.84 ATOM 5288 CG2 VAL B 285 65.037 -11.470 71.478 1.00 26.93 ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.38 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD1 PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5298 CE2 PHE B 286 59.205 -14.180 70.319 1.00 22.92 ATOM 5298 CE2 PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 22.992 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 22.828 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38				GLU B	284	69.084	-7.281	68.222	1.00 57.37
ATOM 5282 C GLU B 284 64.638 -10.105 67.714 1.00 31.93 ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5284 N VAL B 285 64.089 -10.340 68.901 1.00 28.09 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5287 CG1 VAL B 285 63.257 -13.206 71.262 1.00 29.84 ATOM 5288 CG2 VAL B 285 65.037 -11.470 71.478 1.00 26.93 ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5294 CG PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD1 PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5296 CD2 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5298 CE2 PHE B 236 59.205 -14.180 70.319 1.00 22.51 ATOM 5299 CZ PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 22.92 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 22.92 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38					284	68.150	-5.485	69.069	1.00 55.42
ATOM 5283 O GLU B 284 64.899 -11.001 66.913 1.00 28.26 ATOM 5284 N VAL B 285 64.089 -10.340 68.901 1.00 28.09 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5287 CG1 VAL B 285 63.257 -13.206 71.262 1.00 29.84 ATOM 5288 CG2 VAL B 285 65.037 -11.470 71.478 1.00 26.93 ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5290 O VAL B 285 62.460 -12.265 68.758 1.00 31.38 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5294 CG PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5298 CE2 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 22.92 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 26.75 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38								67.714	1.00 31.93
ATOM 5284 N VAL B 285 64.089 -10.340 68.901 1.00 28.09 ATOM 5285 CA VAL B 285 63.765 -11.697 69.325 1.00 30.67 ATOM 5286 CB VAL B 285 63.687 -11.802 70.863 1.00 28.33 ATOM 5287 CG1 VAL B 285 63.257 -13.206 71.262 1.00 29.84 ATOM 5288 CG2 VAL B 285 65.037 -11.470 71.478 1.00 26.93 ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5290 O VAL B 285 62.422 -13.423 68.349 1.00 31.38 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5294 CG PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD1 PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5296 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5298 CE2 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 22.84 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 22.84 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38									
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ATOM 5289 C VAL B 285 62.460 -12.265 68.758 1.00 31.19 ATOM 5290 O VAL B 285 62.422 -13.423 68.349 1.00 31.38 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5294 CG PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 386 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27			CG2	VAL B	285				
ATOM 5290 O VAL B 285 62.422 -13.423 68.349 1.00 31.38 ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5294 CG PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD1 PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5296 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5298 CE2 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5299 CZ PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 22.84 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38				VAL B	285	62.460	-12.265		
ATOM 5291 N PHE B 286 61.398 -11.460 68.729 1.00 28.21 ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5294 CG PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.92 ATOM 5298 CE2 PHE B 286 59.880 -13.213 72.833 1.00 22.92 ATOM 5299 CZ PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38								68.349	1.00 31.38
ATOM 5292 CA PHE B 286 60.105 -11.948 68.249 1.00 25.71 ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5294 CG PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5295 CD1 PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD2 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5298 CE2 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.388									1.00 28.21
ATOM 5293 CB PHE B 286 59.064 -11.853 69.374 1.00 24.57 ATOM 5294 CG PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5298 CE2 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5299 CZ PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 59.518 -11.318 66.930 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 23.38	-	-							
ATOM 5294 CG PHE B 286 59.311 -12.804 70.514 1.00 26.87 ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 386 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 65.130 1.00 23.38				rns s	200				
ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 386 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27	MOTA			PHE B	200				
ATOM 5295 CD1 PHE B 286 59.651 -12.331 71.779 1.00 25.16 ATOM 5296 CD2 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5299 CZ PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27	ATOM	5294	CG	PHE B	286				
ATOM 5296 CD2 PHE B 286 59.205 -14.180 70.319 1.00 22.51 ATOM 5297 CE1 PHE B 286 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 286 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27		5295	CD1	PHE B	286			/1.//9	
ATOM 5297 CE1 PHE B 236 59.880 -13.213 72.833 1.00 22.92 ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5300 C PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5301 O PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27			CD2	PHE B	286				
ATOM 5298 CE2 PHE B 236 59.433 -15.063 71.362 1.00 21.99 ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 22.84 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27			CE1	PHE B	286	59.880	-13.213		
ATOM 5299 CZ PHE B 286 59.772 -14.578 72.626 1.00 26.75 ATOM 5300 C PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5301 O PHE B 286 58.388 -11.630 66.620 1.00 22.84 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27			CE2	PHF A	236			71.362	1.00 21.99
ATOM 5299 C2 PHE B 286 59.518 -11.318 66.993 1.00 25.90 ATOM 5300 C PHE B 286 58.388 -11.630 66.620 1.00 22.84 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27				DHE P	286				1.00 26.75
ATON 5300 C FALL B 286 S8.388 -11.630 66.620 1.00 22.84 ATOM 5302 N GLY B 287 60.272 -10.451 66.329 1.00 28.27 ATOM 5302 N GLY B 287 60.272 -10.451 65.130 1.00 23.38				DAE D	286				
ATOM 5300 N GLY B 287 60.272 -10.451 66.329 1.00 28.27				PAGE D	206				
ATOM 3302 C1 C11 T 297 50 756 -9 814 65.130 1.00 23.38	MOTA.			PHE B	200				
	MOTA			GLY B	28/				1 00 23 39
		5303	CA	GLY B	287	59.756	-A.8T4	- 05.250	1.00 23.30

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1.00 29.17
                                    58.765
                                            -8.719
                                                      65.498
                   GLY B 287
ATCM
        5304
              С
                                    58.786
                                                      56.617
                                                              1.00
                                                                    22.88
                                             -8.216
        5305
                   GLY B 287
ATCM
                                                              1.00 26.77
                                                      64.558
                                             -8.361
        5306
                   GLU B 288
                                    57.896
              N
ATOM
                                             -7.324
                                                      64.754
                                                              1.00 25.38
                                    56.893
                   GLU B 288
ATOM
        5307
              CA
                                                      63.405
                                                               1.00
                                    56.405
                                             -6.791
                                                                    29.51
                   GLU B 288
        5308
MOTA
                                                              1.00 36.06
                                             -6.003
                                                      62.605
                                    57.430
               CG
                   GLU B 288
        5309
ATCM-
                                                               1.00 41.10
                                    57.906
                                             -4.769
                                                      63.347
        5310
               CD
                   GLU B
                         288
ATOM
                                             -4.055
                                                               1.00 41.19
                                    57.058
                                                      63.919
                   GLU B 288
        5311
               OE1
ATCM
                                                               1.00 44.69
                                             -4.503
                                                      63.348
                   GLU B 288
                                    59.125
               OE2
MOTA
        5312
                                                               1.00 27.87
                                             -7.819
                                                      65.527
                   GLU B 288
                                    55.682
        5313
               С
ATOM
                                   55.209
                                             -8.931
                                                               1.00 26.80
                                                      65.308
        5314
                   GLU B 288
MOTA
                                                               1.00 24.53
                                             -6.973
                                                      66.419
                                    55.176
                   GLY B 289
MOTA
        5315
              N
                                    54.006
                                             -7.326
                                                      67.204
                                                               1.00 29.17
                   GLY B 289
        5316
              CA
ATOM
                                    53.015
                                                               1.00 30.46
                                             -6.171
                                                      67.244
                   GLY B 289
        5317
ATOM
                                             -5.326
                                                      66.358
                                                               1.00 26.17
                   GLY B 289
                                    53.005
              0
ATOM
        5318
                                                               1.00 23.95
                   VAL B 290
                                    52.171
                                             -6.142
                                                      68.268
        5319
              N
ATOM
                                    51.194
                                                      68.440
                                                               1.00 22.25
                                             -5.079
               CA
                   VAL B 290
        5320
ATOM
                                    49.794
                                             -5.655
                                                      68.783
                                                               1.00 18.71
                   VAL B 290
               CB
        5321
ATOM
                                                               1.00 22.67
                                                      69.047
                                    48.810
                                             -4.525
                   VAL B 290
              CG1
ATOM
        5322
                                    49.289
                                             -6.504
                                                      67.629
                                                               1.00 19.26
                   VAL B 290
        5323
               CG2
ATOM
                                    51.722
                                             -4.232
                                                      69.593
                                                               1.00 21.55
                   VAL B
                         290
        5324
ATOM
                                             -4.741
                                                      70.687
                                                               1.00 21.32
                                    51.960
                   VAL B 290
        5325
               0
ATOM
                                                               1.00 21.06
                   TYR B 291
                                    51.913
                                             -2.941
                                                      69.346
        5326
              N
MOTA
                                    52.479
                                             -2.063
                                                      70.357
                                                               1.00 19.29
ATOM
        5327
               CA
                   TYR B
                         291
                                    53.582
                                             -1.216
                                                      69.711
                                                               1.00 20.40
                   TYR B 291
        5328
               CB
ATOM
                                                               1.00 23.09
                                             -2.072
                                                      68.918
                                    54.553
                   TYR B 291
               CG
MOTA
        5329
                                                               1.00 19.52
                   TYR B 291
                                    54.740
                                             -1.875
                                                      67.549
        5330
              CD1
ATOM
                                    55.580
                                                      66.809
                                             -2.712
                                                               1.00 20.67
                   TYR B 291
ATOM
        5331
               CE1
                                                               1.00 22.88
                                                      69.527
                                    55.234
                                             -3.122
                   TYR B 291
        5332
               CD2
ATOM
                                                               1.00 26.04
                                                      68.800
                   TYR B 291
                                    56.070
                                             -3.960
               CE<sub>2</sub>
        5333
MOTA
                                    56.235
                                             -3.752
                                                      67.442
                                                               1.00 23.44
                   TYR B 291
               CZ
MOTA
        5334
                                    57.027
                                                      66.722
                                                               1.00 28.02
                                             -4.612
                   TYR B
                         291
        5335
               OH
MOTA
                                                      71.068
                                             -1.180
                                                               1.00 26.89
                   TYR B 291
                                    51.465
ATOM
        5336
               C
                                                               1.00 20.26
                   TYR B 291
                                    50.668
                                             -0.479
                                                      70.429
        5337
ATOM
                                             -1.204
                                                      72.399
                                                               1.00 21.75
                                    51.522
ATOM
        5338
               N
                   LEU B
                         292
                                                      73.227
                                                               1.00 22.11
                                    50.604
                                             -0.426
                   LEU B 292
        5339
               CA
MOTA
                                                      74.088
                                                               1.00 20.92
                                             -1.369
                         292
                                    49.765
                   LEU B
        5340
               CB
MOTA
                                                      73.375
                                                               1.00 22.94
                                    49.091
                                             -2.542
                   LEU B 292
ATOM
        5341
               CG
                                                               1.00 21.03
                                             -3.362
                                                      74.411
                                    48.328
        5342
               CD1
                   LEU B 292
ATOM
                                    48.149
                                             -2.043
                                                      72.281
                                                               1.00 18.04
                         292
               CD2
                   LEU B
        5343
ATOM
                                                               1.00 21.59
                                                      74.147
                   LEU B 292
                                    51.330
                                              0.557
MOTA
        5344
                                                               1.00 19.96
                                                      74.426
                                    52.514
                                              0.404
                   LEU B 292
MOTA
        5345
               0
                                                               1.00 23.31
                                              1.571
                                                      74.613
                                    50.606
              N
                   GLY B 293
        5346
ATOM
                                    51.195
                                                               1.00 20.76
                                                      75.521
                   GLY B 293
                                              2.537
              CA
        5347
MOTA
                                    51.163
                                              1.979
                                                      76.930
                                                               1,00 26.15
                   GLY B 293
        5348
               C
ATOM
                                                               1.00 20.96
                                    51.263
                                              0.765
                                                      77.133
        5349
                   GLY B 293
ATOM
                                                      77.914
                                                               1.00 24.63
                                    51.017
                                              2.859
                   GLY B 294
        5350
              N
ATOM
                                                               1.00 20.00
                                              2.407
                                                      79.293
                   GLY B 294
                                    50.980
              CA
        5351
ATOM
                                                               1.00 22.59
                                                      80.285
                                    51.176
                                              3.538
                   GLY B 294
ATOM
        5352
              С
                                                               1.00 17.46
                                    51.145
                                              4.719
                                                      79.916
                   GLY B 294
        5353
ATOM
                                                               1.00 17.10
                                                      81.551
                                    51.373
                                              3.179
        5354
                   GLY B 295
              N
ATOM
                                                               1.00 16.52
                                                      82.582
                   GLY B 295
                                    51.577
                                              4.180
        5355
              CA
MCTA
                                                               1.00 19.54
                                    52.695
                                              5.145
                                                      82.232
                   GLY B 295
        5356
ATOM
                                              4.737
                                                      81.732
                                                               1.00 16.31
                                    53.738
                         295
              0
                   GLY B
ATOM
        5357
                                                      82.497
                                                               1.00 21.93
                   GLY B 296
                                    52.467
                                              6.430
        5358
              N
ATCM
                                                               1.00 20.05
                                    53.448
                                              7.465
                                                      82.207
                   GLY B 296
        5359
               CA
ATCM
                                    52.869
                                              8.750
                                                      82.759
                                                               1.00 22.20
                         296
                   GLY B
        5360
ATOM
                                                      82.336
                                                               1.00 20.48
                                    51.790
                   GLY B 296
                                              9.160
        5361
ATOM
                                                               1.00 20.93
                                    53.573
                                              9.402
                                                      83.682
                   TYR B 297
        5362
               N
ATOM
                                                               1.00 23.25
                                    53.025
                                             10.598
                                                      84.306
                         297
               CA
                   TYR B
        5363
ATOM
                                                      85.774
                                                               1.00 19.93
                                             10.284
                   TYR B 297
                                    52.731
        5364
               CB
ATOM
                                                               1.00 24.76
                                              8.944
                                                      85.900
                   TYR B 297
                                    52.041
        5365
               CG
ATOM
                                                               1.00 21.97
                                    52.779
                                              7.758
                                                      85.936
                          297
               CD1
                   TYR B
        5366
ACOM
                                                               1.00 19.79
                                                      85.912
               CE1 TYR B 297
                                    52.148
                                              6.514
ATOM
        5367
                                                               1.00 20.86
                                                      85.849
               CD2 TYR B 297
                                    50.653
                                              8.850
        5368
                                                               1.00 19.57
ATOM
                                    50.012
                                               7.612
                                                      85.822
               CE2
                   TYR B
                         297
ATOM
        5369
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Figure 18-82

ATOM	5370 CZ TYR B 297				
		, 50.758			1.00 23.85
ATOM	5371 OH TYR B 297	50.106	5 .5.254	85.806	1.00 17.83
MOTA	5372 C TYR B 297	53.839		84.181	
MOTA		53.451	12.925	84.705	1.00 21.77
MOTA	5374 N HIS B 298	54.974	11.794	83.497	
ATOM	5375 CA HIS B 298	55.787			
				83.270	
MOTA	5376 CB HIS B 298	57.270	12.713	83.534	1.00 22.88
MOTA	5377 CG HIS B 298	58.097	13.956	83.502	1 00 35 13
ATOM	5378 CD2 HIS B 298				
		58.406		82.482	1.00 28.22
MOTA	5379 ND1 HIS B 298	58.617	14.536	84.641	1.00 32.76
MOTA	5380 CE1 HIS B 298	59.209			1.00 32.70
				84.323	1.00 26.52
ATOM	5381 NE2 HIS B 298	59.094	15.852	83.019	1.00 32.15
ATOM	5382 C HIS B 298	55.589	13.307	81.795	1.00 25.66
ATOM	5383 O HIS B 298				
		56.087		80.923	1.00 25.84
MOTA	5384 N PRO B 299	54.901	14.424	81.496	1.00 27.02
ATOM	5385 CD PRO B 299	54.388			
				82.447	1.00 29.91
ATOM	5386 CA PRO B 299	54.616	14.864	80.127	1.00 26.53
ATOM	5387 CB PRO B 299	53.952	16.232	80.342	1.00 27.76
MOTA	5388 CG PRO B 299	54.583			
			16.696	81.656	1.00 27.97
MOTA	5389 C PRO B 299	55.815	14.930	79.194	1.00 27.08
ATOM	5390 O PROB 299	55.738	14.472	78.057	1.00 28.58
MOTA	5391 N TYR B 300				
		56.925	15.484	79.668	1.00 27.30
ATOM	5392 CA TYR B 300	58.114	15.593	78.824	1.00 27.17
MOTA	5393 CB TYR B 300	59.173	16.496	79.466	
					1.00 31.65
MOTA		58.684	17.851	79.921	1.00 31.61
MOTA	5395 CD1 TYR B 300	57.414	18.318	79.582	1.00 32.71
ATOM	5396 CE1 TYR B 300 ·	56.971			
			19.568	80.014	1.00 38.52
ATOM	5397 CD2 TYR B 300	59.499	18.670	80.701	1.00 30.92
ATOM	5398 CE2 TYR B 300	59.072	19.517	·81.138	1.00 32.13
MOTA	5399 CZ TYR B 300				
	·	57.808	20.361	80.795	1.00 39.17
MOTA	5400 OH TYR B 300	57.374	21.585	81.252	1.00 43.90
MOTA	5401 C TYR B 300	58.731	14.218	78.572	1.00 25.20
ATOM					
		59.106	13.894	77.445	1.00 25.15
MOTA	5403 N ALA B 301	58.845	13.419	79.628	1.00 20.55
ATOM	5404 CA ALA B 301	59.414	12.080	79.508	
					1.00 22.12
MOTA	5405 CB ALA B 301	59.417	11.388	80.874	1.00 17.09
ATOM	5406 C ALA B 301	58.608	11.260	78.505	1.00 15.20
MOTA	5407 O ALA B 301	59.161			
			10.629	77.613	1.00 17.12
ATOM	5408 N LEU B 302	57.295	11.290	78.667	1.00 18.02
ATOM	5409 CA LEU B 302	56.381	10.553	77.815	1.00 19.88
ATOM	5410 CB LEU B 302				
		54.957	10.702	78.362	1.00 21.72
ATOM ·	5411 CG LEU B 302	53.767	10.118	77.606	1.00 31.08
ATOM	5412 CD1 LEU B 302	52.576	9.980	78.549	1.00 31.35
ATOM					
		53.434	11.011	76.415	1.00 27.11
ATOM	5414 C LEU B 302	56.445	10.988	76.351	1.00 21.13
ATOM	5415 O LEU B 302	56.473	10.149	75.449	1.00 21.76
ATOM	5416 N ALA B 303				
		56.472	12.293	76.115	1.00 17.69
ATOM	5417 CA ALA B 303	56.516	12.811	74.755	1.00 17.79
ATOM	5418 CB ALA B 303	56.357	14.326	74.780	
					1.00 24.50
ATCM	5419 C ALA B 303	57.803	12.425	74.040	1.00 20.84
ATOM	5420 O ALA B 303	57.781	11.968	72.891	1,00 19.33
	5421 N ARG B 304				1,00 15.55
MOTA		58.930		74.723	1.00 21.08
MOTA	5422 CA ARG B 304	60.215	12.269	74.120	1.00 25.56
ATOM	5423 CB ARG B 304	61.375		74.962	1 00 10 37
					1.00 18.37
ATOM	5424 CG ARG B 304	61.427		75.072	1.00 23.12
ATOM	5425 CD ARG B 304	62.797	14.758	75.624	1.00 29.00
ATOM	5426 NE ARG B 304				
		63.073		76.789	1.00 33.28
MOTA	5427 CZ ARG B 304	64.271	13.689	77.283	1.00 30.24
MOTA	5428 NH1 ARG B 304	65.363			1.00 24.98
				70 70	
ATOM		64.365	12.896	78.333	1.00 36.15
ATOM	5430 C ARG B 304	60.406	10.775		1.00 20.46
ATOM	5431 O ARG B 304	60.850			
					1.00 18.70
MOTA	5432 N ALA B 305	60.070	9.988	74.937	1.00 22.48
ATOM .	5433 CA ALA B 305	60.226			1.00 19.70
ATOM	5434 CB ALA B 305	59.847			
					1.00 24.24
ATOM	5435 C ALA B 305	59.407	7.930	73.711	1.00 15.82
			•		

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MOTA	5436	0	ALA	В	305	59.938	7.184	72.888	1.00 19.12
MOTA	5437	N	TRP	В	306	58.113	8.230	73.659	1.00 18.65
ATOM	5438	CA	TRP	В	306	57.298	7.668	72.600	1.00 19.57
MOTA	5439	CB	TRP	В	306	55.800	7.856	72.893	1.00 18.26
ATOM	5440	CG	TRP	В	306	55.301	6.911	73.953	1.00 20.71
					306	54.087	7.025	74.708	1.00 23.94
MOTA	5441	CD2	TRP						
MOTA	5442	CE2	TRP	В	306	53.988	5.870	75.513	1.00 24.73
	5443	CE3	TRP	ъ	306	53.073	7.991	74.780	1.00 26.01
ATOM									
MOTA	5444	CD1	TRP	В	306	55.872	5.721	74.326	1.00 20.04
ATOM	5445	NE1	TRP	В	306	55.092	5.093	75.260	1.00 19.17
ATOM	5446	CZ2	TRP	B	306	52.912	5.655	76.385	
MOTA	5447	CZ3	TRP	В	306	52.001	7.779	75.646	1.00 28.68
		CH2	TRP		306	51.930	6.619	76.437	1.00 31.22
MOTA	5448								
MOTA	5449	С	TRP	В	306	57.665	8.223	71.226	1.00 23.48
MOTA	5450	0	TRP	В	306	57.416	7.574	70.212	1.00 22.38
									1.00 22.36
ATOM	5451	N	THR	Ħ	307	58.262	9.412	71.176	
ATOM	5452	CA	THR	В	307	58.672	9.953	69.880	1.00 25.94
			THR		307	59.143	11.417	69.986	1.00 25.88
MOTA	5453	CB							
ATOM	5454	OG1	THR	В	307	58.015	12.261	70.258	1.00 21.07
ATOM	5455	CG2	THR	В	307	59.827	11.864	68.686	1.00 22.52
					307	59.815	9.078	69.350	1.00 30.09
MOTA	5456	С	THR						
MOTA	5457	0	THR	В	307	59.922	8.834	68.144	1.00 25.82
MOTA	5458	N	LEU	В	308	60.664	8.596	70.258	1.00 27.54
						61.773	7.734	69.857	1.00 26.76
ATOM	5459	CA	LEU		308				
ATOM	5460	CB	LEU	В	308	62.691	7.424	71.054	1.00 24.24
ATOM	5461	CG	LEU	В	308	63.420	8.614	71.718	1.00 31.16
								72.877	1.00 24.71
MOTA	5462		LEU		308	64.282	8.147		
ATOM	5463	CD2	LEU	В	308	64.289	9.325	70.700	1.00 24.59
ATOM	5464	С	LEU	R	308	61.184	6.443	69.287	1.00 27.20
			LEU		308	61.609	5.961	68.234	1.00 23.52
ATOM	5465	0							
ATOM	546 6	N	ILE	В	309	60.190	5.898	69.980	1.00 25.10
MOTA	5467	CA	ILE	В	309	59.537	4.679	69.530	1.00 25.14
	5468	CB	ILE	В	309	58.387	4.266	70.485	1.00 27.05
ATOM									
MOTA	5469	CG2	ILE	В	309	57.646	3.058	69.926	1.00 23.57
ATOM	5470	CG1	ILE	В	309	58.952	3.94 7	71.868	1.00 22.98
ATOM	5471	CD1	ILE	B	309	59.927	2.793	71.868	1.00 24.25
						58.958	4.885	68.133	1.00 25.41
ATOM	5472	С	ILE						
MOTA	5473	0	ILE	В	309	59.177	4.064	67.243	1.00 22.13
ATOM	5474	N	TRP	В	310	58.232	5.984	67.943	1.00 27.45
						57.618	6.266	66.548	1.00 29.27
MOTA	5475	CA	TRP		310				
ATOM	5476	CB	TRP	В	310	56.721	7.505	66.715	1.00 27.00
ATOM	5477	CG	TRP	В	310	56.112	7.847	65.378	1.00 28.26
						55.172	7.061	64.633	1.00 27.50
ATOM	5478	CD2			310				
ATOM	5479	CE2	TRP	В	310	54.947	7.729	63.408	1.00 30.47
ATOM	5480	CE3	TRP	В	310	54.500	5.856	64.877	1.00 29.85
						56.406	P.929	64.597	1.00 29.76
MOTA	5481	CD1	TRP						
MOTA	5482	NE1	TRP	В	310	55.713	865	63.415	1.00 26.71
ATOM	5483	CZ2	TRP	В	310	54.076	234	62.429	1.00 28.23
						53.636	5.362	63.901	1.00 30.24
ATOM	5484	CZ3	TRP		310				
ATOM	5485	CH2	TRP	В	310	53.433	6.053	62.692	1.00 27.63
ATOM	5486	С	TRP	В	310	58.629	6.424	65.520	1.00 30.16
						58.378	5.964	64.410	1.00 30.04
ATOM	5487	0	TRP						
ATOM	5488	N	CYS	В	311	59.762	7.069	65.793	1.00 24.26
	5489	CA	CYS	R	311	60.782	7.233	64.764	1.00 27.97
ATOM						61.893	8.157	65.252	1.00 28.21
ATOM	5490	CB	CYS						
ATOM	5491	SG	CYS			61.422	9.905	65.381	1.00 33.38
MOTA	5492	С	CYS			61.380	5.886	64.351	1.00 30.02
						61.670	5.660	63.172	1.00 25.45
MOTA	5493	0	CYS		211				
MOTA	5494	N	GLU			61.570	5.001	65.327	1.00 31.59
ATOM	5495	CA	GLU			62.111	3.669	65.067	1.00 33.48
			GLU	2	312	62.142	2.843	66.352	1.00 34.78
ATOM	5496	CB							
MOTA	5497	CG	GLU	Ħ	312	63.487	2.307	66.758	1.00 39.45
MOTA	5498	CD	GLU	В	312	64.171	1.513	65.675	1.00 40.11
	5499	OFI	GLU	В	312	63.539	0.614	65.081	1.00 43.69
ATOM		053	2110	5	212		1.782	65.437	1.00 39.26
ATCM	5500	OE2	GLU	B	214	65.358	1./02		
ATCM	5501	С	GLU	В	312	61.197	2.959	64.080	1.00 29.97

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MOTA	5502	0	.GLU B 312		61.640	2.497	63.035	1.00	31.38
ATOM	5503		LEU B 313		59.919	2.865	64.438		26.70
MOTA	5504	CA			58.930	2.203	63.598		26.73
MOTA	5505	CB	LEU B 313		57.571	2.173	64.297	1.00	25.83
ATOM	5506	CG	LEU B 313		57.429	1.224	65.477	1.00	35.18
ATOM	5507		1 LEU B 313		56.063	1.434	66.130		32.49
MOTA	55Q8	CD	2_LEU B 313		57.595	-0.215	64.989	1.00	29.71
ATOM	5509	С	LEU B 313		58.768	2.866	62.248	1.00	29.03
ATOM	5510	0	LEU B 313		58.716	2.187	61.228	1.00	25.39
MOTA	5511	N	SER B 314		58.677	4.194	62.263		30.13
MOTA	5512	CA	SER B 314	•	58.498	5.006	61.060		34.06
ATOM	5513	CB	SER B 314		58.206	6.456	61.445	1.00	31.15
ATOM	5514	OG	SER B 314		57.041	6.537	62.234	1.00	48.58
ATOM	5515	C	SER B 314		59.707	5.003	60.151		31.84
MOTA	5516	0	SER B 314		59.632	5.469	59.026		34.15
ATOM	5517	N	GLY B 315		60.831	4.515	60.65 5	1.00	31.81
ATOM	5518	CA	GLY B 315		62.036	4.485	59.848	1.00	37.27
MOTA	5519	С	GLY B 315		62.659	5.851	59.616	1.00	39.93
	5520	ŏ	GLY B 315		63.363	6.054	58.624		39.79
MOTA									
ATOM	5521	N	ARG B 316		62.422	6.798	60.518		38.22
MOTA	5522	CA	ARG B 316		63.004	8.121	60.336	1.00	38.66
ATOM	5523	CB	ARG B 316		61.908	9.184	60.275	1.00	40.20
ATOM	5524	CG	ARG B 316		61.089	9.345	61.520		39.00
						10.398			
MOTA	5525	CD	ARG B 316	•	60.032		61.284		42.13
MOTA	5526	NE	ARG B 316		59.002	9.954	60.352		45.09
ATOM	5527	CZ	ARG B 316		58.075	10.754	59.838	1.00	40.84
MOTA	5528	NHI	ARG B 316		58.064	12.033	60.170		48.44
	5529	NH2			57.150	10.278	59.014		35.96
ATOM				•					
ATOM	5530	C	ARG B 316		64.031	8.467	61.408		39.03
ATOM	5531	0	ARG B 316		63.952	7.988	62.539	1.00	34.34
ATOM	5532	N	GLU-B 317		65.003	9.296	61.035	1.00	39.58
ATOM	5533	CA	GLU B 317		66.074	9.697	61.943	1.00	43.35
	5534	СВ	GLU B 317		67.142	10.509	61.203		49.34
MOTA									
ATOM	5535	ĊG	GLU B 317		67.609	9.910	59.884		57.04
ATOM	5536	CD	GLU B 317		66.546	10.009	58.798	1.00	62.79
ATOM	5537	OE1	GLU B 317		66.146	11.149	58.467	1.00	63.46
ATOM	5538	OE2			66.108	8.954	58.280		64.46
	5539	c	GLU B 317		65.555	10.528	63.100		41.58
MOTA									
MOTA	5540	0	GLU B 317		64.658	11.356	62.939		39.74
ATOM .	5541	N	VAL B 318		66.118	10.301	64.278	1.00	35.38
ATOM	5542	CA	VAL B 318		65.706	11.049	65.448	1.00	38.76
ATOM	5543	CB	VAL B 318		66.000	10.265	66.750	1.00	42.28
ATOM	5544	CG1			65.560	11.080	67.962		38.26
MOTA	5545	CG2			65.287	8.916	66.722		39.99
MOTA	5546	С	VAL B 318		66.459	12.370	65.478	1.00	41.82
.TOM	5547	0	VAL B 318		67.689	12.395.	65.570	1.00	37.20
ATOM	5548	N	PRO B 319		65.735	13.491	65.356	1.00	43.18
	5549	CD	PRO B 319		64.290	13.672	65.155		41.90
MOTA									
ATOM	5550	CA	PRO B 319		66.402	14.792	65.388		44.31
MOTA	5551	CB	PRO B 319		65.241	15.763	65.181	1.00	44.58
ATOM	5552	CG	PRO B 319		64.079	15.011	65.795	1.00	43.34
ATOM	5553	C	PRO B 319		67.086	14.965	66.741	1.00	44.62
	5554	ō	PRO B 319		66.541	14.565	67.771		43.75
ATOM									
MOTA	.5555	N	GLU B 320		68.277	15.552	66.735		44.16
MOTA	5556	CY	GLU B 320		69.029	15.762	67.967	1.00	45.92
MOTA	5557	CB	GLU B 320		70.381	16.406	67.663	1.00	50.87
ATOM	5558	CG	GLU B 320		71.165	16.768	68.919		53.71
	5559								
MOTA		CD	GLU B 320		72.455	17.505	68.620		57.75
MOTA	5560		GLU B 320		73.161	17.874	69.583		56.37
MOTA	5561	0E2	GLU B 320		72.762	17.714	67.427	1.00	60.07
ATOM	5562	С	GLU B 320		68.311	16.625	68.995		44.42
	5563	ō	GLU B 320		68.244	16.279	70.168		42.32
ATOM									
MOTA	5564	N	LYS B 321		67.778	17.753	68.550		42.50
MOTA	5565	CA	LYS B 321		67.102	18.672	69.448	1.00	45.24
ATOM	5566	CB	LYS B 321		67.853	20.000	69.503		46.43
	5567	CG	LYS B 321		67.890	20.802	68.195		51.45
MOTA					3330				

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Figure 18-85

> moi/	EE 60	CD	LYS B	321	68.700	20.144	67.057	1.00 57.24
ATOM	5568	CE		321	67.936	19.062	66.280	1.00 55.24
ATOM	5569			321	66.738	19.588	65.558	1.00 55.31
MOTA	5570	NZ		321	65.662	18.971	69.098	1.00 43.44
MOTA	5571	C 0		321	65.211	18.736	67.978	1.00 43.03
MOTA	5572	N		322	64.947	19.512	70.076	1.00 39.45
MOTA	5573			322	63.563	19.885	69.875	1.00 40.31
ATOM	5574	CA		322	62.846	20.034	71.215	1.00 40.88
ATOM	5575	CB		322	62.943	18.901	72.234	1.00 40.09
_ATOM	5576	CG		322	62.001	19.175	73.388	1.00 38.17
ATOM	5577		LEU B		62.588	17.596	71.580	1.00 41:56
ATOM	5578		LEU B		63.615	21.244	69.197	1.00 41.23
MOŢA	5579	C	LEU B		64.466	22.070	69.531	1.00 39.22
MOTA	5580	0	ASN B	323	62.735	21.473	68.233	1.00 40.04
MOTA	5581	N			62.703	22.771	67.582	1.00 43.32
ATOM	5582	CA	ASN B		61.985	22.707	66.234	1.00 41.53
. ATOM	5583	CB CG	ASN B		60.617	22.085	66.335	1.00 41.89
MOTA	5584		ASN B	323	59.889	22.308	67.304	1.00 39.79
ATOM	5585	MD3	ASN B	323	60.243	21.317	65.317	1.00 40.43
ATOM	5586		ASN B		61.949	23.690	68.532	1.00 44.76
MOTA	5587	С 0		323	61.402	23.237	69.539	1.00 45.80
ATOM	5588	N		324	61.902	24.973	68.210	1.00 46.85
MOTA	5589	CA	ASN B		61.234	25.930	69.076	1.00 47.60
ATOM	5590	CB	ASN B		61.460	27.348	68.549	1.00 50.87
ATOM	5591	CG	ASN B		61.089	28.407	69.562	1.00 55.06
ATOM	5592		ASN B		59.925	28.565	69.919	1.00 60.68
ATOM	5593		ASN B		62.091	29.131	70.048	1.00 59.17
TOM	5594	C	ASN B		59.740	25.664	69.249	1.00 43.97
ATOM	5595	0	ASN B		59.190	25.898	70.322	1.00 41.33
MOTA	5596 5597	N	LYS B		59.087	25.168	68.201	1.00 43.49
ATOM	5597 5598	CA		325	57.655	24.892	68.264	1.00 45.95
ATOM	5599	CB	LYS B	325	57.112	24.415	66.909	1.00 48.97
ATOM	5600	CG	LYS B	325	57.212	25.400	65.731	1.00 53.41
ATOM	5601	CD	LYS B		58.582	25.386	65.024	1.00 58.77
MOTA	5602	CE	LYS B		59.700	26.013	65.846	1.00 58.10
MOTA	5603	NZ	LYS B		61.024	25.906	65.178	1.00 53.38
MOTA MOTA	5604	C	LYS B		57.368	23.822	69.309	1.00 45.79
MOTA	5605	ō	LYS B		56.375	23.891	70.034	1.00 43.91
MOTA	5606	N		326	58.245	22.829	69.381	1.00 44.28
ATOM	5607	CA		326	58.078	21.746	70.336	1.00 44.25
ATOM	5608	СВ		326	59.013	20.589	69.986	1.00 41.44
ATOM	5609	С	ALA B	326	58.342	22.233	71.757	1.00 40.92
ATOM	5610	0	ALA B	326	57.639	21.843	72.688	1.00 39.02
ATOM	5611	N	LYS B	327	59.352	23.085	71.922	1.00 38.14
ATOM	5612	CA	LYS B	327	59.689	23.603	73.246	1.00 40.11
ATOM	5613	CB	LYS B	327	60.892	24.552	73.178	1.00 42.36
ATOM	5614	CG	LYS B	327	62.174	23.922	72.(59	1.00 45.78
ATOM	5615	CD		327	63.325	24.926	72.675	1.00 48.46 1.00 49.62
ATOM	5616	CE	LYS B	327	64.594	24.367	72.031	
ATOM	5617	NZ .	LYS B	327	65.108	23.139	72.700	1.00 48.53 1.00 39.17
MOTA	5618	С	LYS B	327	58.500	24.338	73.841	1.00 39.17
ATOM	5619	o	LYS B		58.132	24.112	74.994	1.00 38.87
ATOM	5620	N	GLU B		57.898	25.215	73.048	1.00 41.06
ATOM	5621	CA	GLU B		56.750	25.986	73.512	1.00 44.02
ATOM	5622	CB	GLU B		56.357	27.028	72.463 72.258	1.00 44.02
ATOM	5623	CG	GLU B		57.434	28.084		1.00 44.80
TOM	5624	CD	GLU B		57.835	28.742	73.569 74.237	1.00 48.40
ATOM	5625		GLU B		56.949	29.317		1.00 51.20
ATOM	5626	OE2	GLU B		59.029	28.680	73.935	1.00 47.81
ATOM	5627	С	GLU B		55.569	25.087	73.839	
ATOM	5628	0	GLU B		54.794	25.377	74.750	1.00 41.20 1.00 35.31
ATOM	5629	N	LEU B		55.429	23.999	73.090	1.00 32.69
ATOM	5630	CA	LEU B		54.349	23.056	73.334	1.00 32.09
ATCM.	5631	CB	LEU B		54.404	21.900	72.334	1.00 35.00
ATOM	5632	CG	LEU B		53.344	20.813	72.544	1.00 35.01
ATOM	5633	CD1	LEU B	329	51.958	21.430	.72.419	1.00 30.50
•••						F 001		

MOTA	5634	CD2 LEU	3 3 2 0	53.521	19.699	71.525	1.00 32.36
ATOM	5635			54.504	22.507	74.747	1.00 32.36
MOTA	5636			53.621	22.664	75.583	
ATOM	5637			55.640		75.013	
ATOM	5638			55.889			1.00 32.74
ATOM	5639			57.267	21.311	76.330	1.00 34.99
					20.642	76.382	1.00 37.01
MOTA	5640			57.466	19.428	75.470	1.00 34.91
ATOM	5641			58.832	18.817	75.728	1.00 34.69
ATOM	5642			56.369	18.396	75.742	1.00 34.10
MOTA	5643	C LEU E		55.789	22.363	77.429	1.00 37.12
MOTA	5644	O LEU E		55.210	22.110	78.482	1.00 34.19
ATOM	5645	N LYS E		56.353	23.540	77.186	1.00 34.34
MOTA	5646	CA LYS E		56.313	24.604	78.181	1.00 43.35
MOTA	5647	CB LYS B		57.162	25.788	77.712	1.00 46.25
MOTA	5648	CG LYS B		58.658	25.496	77.685	1.00 51.07
ATOM	5649	CD LYS B		59.482	26.610	77.021	1.00 49.96
ATOM	5650	CE LYS B		59.371	27.957	77.733	1.00 53.08
ATOM	5651	NZ LYS B		58.013	28.569	77.662	1.00 56.18
MOTA	5652	C LYS B		54.892	25.069	78.494	1.00 42.06
MOTA	5653	O. LYS B		54.588	25.416	79.631	1.00 43.05
ATOM	5654	N SER B		54.018	25.056	77.492	1.00 44.54
ATOM	5655	CA SER B		52.639	25.502	77.679	1.00 46.58
ATOM	5656	CB SER B		51.975	25.751	76.329	1.00 48.75
ATOM	5657	OG SER B	332	51.769	24.527	75.646	1.00 49.55
MOTA	5658	C SER B		51.780	24.507	78.451	1.00 49.56
MOTA	5659	O SER B	332	50.618	24.791	78.749	1.00 46.67
MOTA	5660	N ILE B	333 .	52.341	23.345	78.770	1.00 50.55
MOTA	5661	CA ILE B	333	51.586	22.326	79.488	1.00 51.93
MOTA	5662	CB ILE B	333	52.259	20.945	79.376	1.00 51.82
MOTA	5663	CG2 ILE B	333	51.447	19.902	80.134	1.00 50.29
ATOM	5664	CG1 ILE B	333	52.359	20.539	77.905	1.00 52.18
ATOM	5665	CD1 ILE B	333	53.044	19.210	77.693	1.00 55.42
ATOM	5666		333	51.367	22.634	80.964	1.00 51.45
ATOM	5667	O IĻE B	333 .	52.180	23.290	81.614	1.00 50.96
ATOM	5668	N ASP B	334	50.245	22.141	81.472	1.00 54.05
ATOM	5669	CA ASP B		49.850	22.306	82.865	1.00 58.15
MOTA	5670	CB ASP B		48.320	22.216	82.959	1.00 60.38
ATOM	. 5671	CG ASP B		47.751	20.972	82.262	1.00 63.85
MOTA	5672	OD1 ASP B		48.017	19.833	82.710	1.00 59.16
ATOM	5673	OD2 ASP B	334	47.033	21.138	81.252	1.00 59.71
ATOM	5674	C ASP B		50.506	21.207	83.701	1.00 55.47
MOTA	5675	O ASP B		49.833	20.291	84.171	1.00 54.08
ATOM	5676		335	51.816	21.307	83.906	1.00 54.60
MOTA	5677		335	52.524	20.266	84.641	1.00 56.60
ATOM	5678	CB PHE B	335	53.718	19.784	83.811	1.00 53.01
MOTA	5679	CG PHE B		54.522	18.717	84.482	1.00 49.30
MOTA	5686	CD1 PHE B		53.898	17.589	85.008	1.00 45.61
ATOM	5681	CD2 PHE B		55.901	18.843	84.605	1.00 46.83
MOTA	5682	CE1 PHE B		54.637	16.600	85.651	1.00 45.95
MOTA	5683	CE2 PHE B		56.651	17.860	85.247	1.00 46.02
MOTA	5684	CZ PHE B		56.018	16.737	85.772	1.00 46.08
MOTA	5685	C PHE B		52.971	20.559	86.072	1.00 57.29
ATOM	5686	O PHE B		52.197	20.378	87.012	1.00 63.54
MOTA	5687	N GLU B	336	54.223	20.983	86.229	1.00 55.21
MOTA	5688	CA GLU B	336	54.818	21.286	87.535	1.00 60.30
MOTA	5689	CB GLU B	336	53.783	21.846	88.517	1.00 64.95
MOTA	5690	CG GLU B		54.375	22.225	89.867	1.00 71.50
ATOM	5691	CD GLU B	336	53.363	22.882	90.787	1.00 75.37
MOTA	5692	OE1 GLU B		52.796	23.925	90.394	1.00 75.32
ATOM	5693	OE2 GLU B	336	53.137	22.361	91.901	1.00 76.84
MOTA	5694	C GLU B		55.485	20.058	88.146	1.00 55.66
ATOM	5695	O GLU B		54.823	19.093	88.529	1.00 49.97
MOTA	5696	N GLU B		56.807	20.125	88.240	1.00 54.26
ATOM ·	5697	CA GLU B		57.630	19.047	88.767	1.00 54.35
ATOM	5698	CB GLU B		59.101	19.457	88.635	1.00 54.08
ATOM	5699	CG GLU B	337	60.074	18.315	88.514	1.00 54.15
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ATOM	5700	CD	GLU B	33 7	59.856	17.496	87.259	1.00 48.94
ATOM	5701	OE1	GLU B	337	59.958	18.049	86.142	1.00 41.06
ATOM	5702	OE2	GLU B		59.581	16.292	87.391	1.00 50.23
				337	57.278	18.740	90.227	1.00 55.08
MOTA	5703	C						1.00 54.51
MOTA	5704	0		337	57.130	19.651	91.039	
ATOM	5705	N	PHE B	338	57.140	17.458	90.557	1.00 56.20
ATOM	5706	CA	PHE B	338	56.798	17.048	91.918	1.00 57.73
MOTA	5707	CB	PHE B		56.713	15.527	92.020	1.00 58.47
			PHE B		56.231	15.034	93,359	1.00 63.17
MOTA	5708	CG						
MOTA	5709	CDI	PHE B		54.882	15.096	93.696	1.00 64.86
ATOM	5710	CD2	PHE B	338	57.129	14.526	94.293	1.00 63.51
ATOM	5711	CE1	PHE B	338	54.434	14.656	94.943	1.00 65.25
ATOM	5712	CE2	PHE B		56.693	14.087	95.539	1.00 63.64
		CZ	PHE B		55.342	14.152	95.864	1.00 66.30
MOTA	5713				57.836	17.539	92.918	1.00 61.85
ATOM	5714	C	PHE B				94.078	
ATOM	5715	0	PHE B		57.520	17.807		1.00 58.15
MOTA	5716	N	ASP B	339	59.081	17.636	92.466	1.00 64.63
ATOM	5717	CA	ASP B	339	60.167	18.099	93.316	1.00 67.53
MOTA	5718	CB	ASP B	339	61.286	17.059	93.362	1.00 67.32
	5719	CG	ASP B		62.474	17.524		1.00 68.13
ATOM						17.909	95.346	1.00 68.68
ATOM	5720	OD1	ASP B		62.280			
ATOM	5721	OD2	ASP B		63.603	17.502	93.646	1.00 69.03
MOTA	5722	С	ASP B	339	60.718	19.435	92.829	1.00 69.03
ATOM	5723	0	ASP B	339	61.211	19.545	91.708	1.00 67.54
ATOM	5724	N	ASP B	340	60.626	20.442	93.693	1.00 72.19
MOTA	5725	CA		340	61.088	21.797	93.402	1.00 75.20
	5726	CB		340	61.113	22.623	94.689	1.00 77.04
ATOM					59.766	22.671	95.375	1.00 78.70
ATOM	5727	CG		340				1.00 79.66
MOTA	5728	OD1		340	58.803	23.181	94.763	
ATOM	5729	OD2	ASP B	340	59.668	22.194	96.525	1.00 80.53
MOTA	5730	C	ASP B	340	62.464	21.856	92.751	1.00 74.82
ATOM	5731	0	ASP B	340	62.615	22.400	91.659	1.00 78.48
ATOM	5732	N		341	63.465	21.303	93.426	1.00 74.11
	5733	CA		341	64.827	21.312	92.907	1.00 76.25
ATOM				341	65.818	21.596	94.040	1.00 79.54
ATOM	5734	CB				21.653	93.596	1.00 82.33
MOTA	5735	CG		341	67.277			
MOTA	5736	CD	GLU B	341	67.539	22.750	92.577	1.00 83.24
ATOM	5737	OE1	GLU B	341	67.333	23.937	92.910	1.00 85.25
ATOM	5738	OE2	GLU B	341	67.950	22.427	91.443	1.00 83.72
ATOM	5739	С	GLU B	341	65.196	19.998	92.227	1.00 73.97
ATOM	5740	ō	GLU B		65.627	19.051	92.883	1.00 77.10
	5741	N		342	65.033	19.946	90.910	1.00 71.92
MOTA				342	65.354	18.744	90.151	1.00 68.51
MOTA	5742	CA					89.663	1.00 68.83
ATOM	5743	CB		342	64.081	18.027		
MOTA	5744			342	63.268	17.552	90.837	1.00 67.57
MOTA	5745	CG2	VAL B	342	63.255	18.969	88.806	1.00 67.72
ATOM	5746	С	VAL B	342	66.201	19.059	88.927	1.00 65 35
ATOM	5747	0	VAL B	342	67.177	18.366	88.640	1.00 68.31
	5748	N	ASP'B		65.819	20.112	88.213	1.00 60.89
MOTA			ASP B		66.514	20.520	86.998	1.00 58.89
ATOM	5749	CA					87.223	1.00 63.48
MOTA	5750	CB	ASP B		68.024	20.636		
ATOM	5751	CG	ASP B	343	68.763	21.070	85.966	1.00 66.69
ATOM	5752	OD1	ASP B	343	70.012	21.070	85.970	1.00 67.64
ATOM	5753		ASP B		68.089	21.420	84.973	1.00 65.42
	5754		ASP B		66.264	19.499	85.900	1.00 53.17
ATOM	5755		ASP B		66.993	18.516	85.766	1.00 49.70
ATOM						19.735	85.124	1.00 50.24
ATOM	5756		ARG B		65.216		84.022	1.00 46.49
MOTA	5757		ARG B		64.868	18.853		1 00 40 43
ATOM	5758	CB	ARG B	344	63.467	18.269	84.228	1.00 42.41
MOTA	5759	CG	ARG B	344	63.317	17.367	85.452	1.00 38.59
ATOM	5760		ARG B		64.344	16.246	85.432	1.00 37.12
			ARG B		64.169	15.310	86.537	1.00 36.55
ATOM	5761		ARG B		65.078	14.413	86.905	1.00 37.20
MOTA	5762					14.331	86.259	1.00 33.53
ATOM .	5763		ARG B		66.234	_	87.915	1.00 28.79
MOTA	5764		ARG B		64.830	13.595		
MOTA	5765	С	ARG B	344	64.910	19.660	82.732	1.00 44.45
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ATOM	5766	0	ARG	B 344		64.328	19.269	81.720	1 00	38.73
ATOM	5767			B 345						
-						65.618		82.783	1.00	42.44
ATOM	5768	CA	SER	B 345		65.740	21.677	81.637	1.00	41.74
ATOM	5769	CE	SER	B 345		66.661	22.849	81.993		43.47
ATOM	5770	OG		B 345						
						67.956		82.351		46.96
MOTA	5771			B 345		66.244	20.981	80.375	1.00	36.32
MOTM	5772	0	SER	B 345		65.840	21.333	79.273	1.00	35.70
ATOM	5773	N	TYR	B 346		67.117	19.992	80.534		33.93
	5774									
MOTA				B 346		67.661	19.264	79.391	1.00	34.77
ATOM	5775		TYR	B 346		68.660	18.206	79.877	1.00	36.09
ATOM	5776	CG	TYR :	B 346	_	68.054	17.146	80.774		34.27
MOTA	5777	CD	1 TYR 1	B 346	-	67.433	16.013	80.240		37.62
MOTA	5778	CE				66.843	15.048	81.077		36.73
ATOM	5779	CD	2 TYR 1	B 346		68.072	17.294	82.157	1.00	36.40
MOTA	5780	CE	2 TYR I	346		67.489	16.344	82.999		36.54
ATOM	5781	CZ	TYR I			66.878	15.228			36.54
•								82.457		
ATOM	5782	ОН				66.310	14.306	83.306	1.00	33.35
ATOM	5783	С	TYR I	3 3 4 6		66.563	18.599	78.570	1.00	36.26
MOTA	5784	0	TYR I	3 3 4 6		.66.719	18.385	77.367		40.50
ATOM	5785	N	MET E							
					٠	65.445	18.282	79.214		32.72
ATOM	5786	CA	MET E			64.346	17.628	78.516	1.00	35.43
MOTA	5787	CB	MET E	3 3 4 7		63.280	17.164	79.513	1.00	34.36
MOTA	5788	CG	MET E	347		63.819	16.292	80.635		28.32
ATOM	5789	SD	MET E							
						62.515	15.604	81.669		34.47
ATOM	5790	CE	MET E			61.654	17.027	82.142	1.00	39.60
MOTA	5791	C	MET E	347		63.701	18.525	77.465		39.04
ATOM	5792	0	MET B			63.060	18.029	76.540		37.38
	5793									
ATOM		N	LEU B			63.857	19.839	77.606		39.21
MOTA	5794	CA	LEU B	348		63.272	20.773	76.645	1.00	40.81
ATOM	5795	CB	LEU B	348		62.806	22.058	77.339	1.00	36.87
ATOM	5796	CG	LEU. B	348		61.690	21.975	78.384		42.66
MOTA	5797		LEU B			61.507	23.337	79.032		43.41
MOTA	5798	CD2	LEU B	348		60.391	21.511	77.741	1.00	40.47
MOTA	5799	С	LEU B	348		64.289	21,133	75.573	1.00	41.13
ATOM	5800	Ō	LEU B			64.018	21.968	74.711		38.93
MOTA	5801	N	GLU B			65.455	20.495	75.632		37.70
ATOM	5802	CA	GLU B	349		66.527	20.757	74.681	1.00	42.48
ATOM	5803	CB	GLU B	349		67.856	20.953	75.422	1.00	45.02
ATOM	5804	CG	GLU B			67.834	22.035	76.493		53.82
MOTA	5805	CD	GLU B			67.483	23.402	75.938		57.46
MOTA	5806	OE1	GLU B	349		68.211	23.885	75.044	1.00	59.62
MOTA	5807	OE2	GLU B	349		66.480	23.993	76.397	1.00	57.91
MOTA	5808	С	GLU B	349		66.709	19.638	73.664		43.57
	5809									
MOTA		0	GLU B	349		66.577	19.849	72.459		41.26
MOTA	5810	N	THR B	350		67.027	18.448	74.161	1.00	41.95
MOTA	5811	CA	THR B	350		67.264	17.299	73.298	1.00	40.02
MOTA	5812	CB		350		68.689	16.775	73.504		43.08
MOTA	5813	OG1		350		68.894	16.490	74.894		41.07
ATOM .	5814	CG2	THR B	350		69.703	17.816	73.049	1.00	45.05
ATOM	5815	С	THR B	350		66.278	16.154	73.510	1.00	37.56
ATOM	5816	ō	THR B			65.754	15.966			33.64
								74.611		
MOTA	5817	N	LEU B			66.043	15.391	72.445		32.86
ATOM	5818	CA	LEU B	351		65.126	14.260	72.475	1.00	35.00
ATOM	5819	CB	LEU B	351		64.776	13.810	71.053		31.61
MOTA	5820	CG	LEU B			63.709	14.601	70.312		35.31
MOTA	5821			351		63.552	14.064	68.904	1.00	37.88
ATOM	5822	CD2	LEU B			62.397	14.474	71.068	1.00	39.36
ATOM	5823	C	LEU B			65.662	13.065	73.240		33.33
ATOM	5824	0	LEU B			64.956	12.469	74.046		31.48
ATOM	5825	N	LYS B	352		66.915	12.720	72.981	1.00	29.58
ATOM	5826	CA		352		67.527	11.576	73.633	1.00	
	5827	СВ		352					1.00	
ATOM						68.457	10.864	72.647.		
ATOM .	5828	CG	LYS B			67.777	10.563	71.326	1.00	19.29
ATOM	5829	CD	LYS B	352		68.703	9.949	70.294	1.00	42.25
ATOM	5830	CE.	LYS B			69.110	8.541	70.655	1.00	46 22
									1 00	44.65
ATOM	5831	NZ	LYS B	J 7 Z		69.831	7.905	69.516	1.00	44.15

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3 moss	5832	C LYS B 352	68.295	11.983	74.878	1.00 36.30
ATOM						
MOTA	5833		69.086	12.931	74.865	1.00 36.65
ATOM	5834	N ASP B 353	68.049	11.275	75.967	1.00 30.01
MOTA	5835	CA ASP B 353	68.757	11.569	77.188	1.00 33.99
ATOM	5836	CB ASP B 353	67.852	11.308	78.394	1.00 38.57
	5837		67.134	9.986	78.315	1.00 43.90
ATOM						
ATOM	5838		66.034	9.851	78.926	1.00 22.39
MOTA	5839		67.679	9.078	77.649	1.00 50.42
ATOM	5840	C ASP B 353	70.022	10.723	77.202	1.00 35.83
ATOM	5841	O _ASP B 353	70.189	9.833	76.368	1.00 23.71
ATOM	5842	N PRO B 354	70.954	11.025	78.116	1.00 36.36
ATOM	5843		70.928	12.093	79.132	1.00 38.28
•	5844		72.205	10.277	78.212	1.00 33.62
ATOM			73.003			1.00 34.46
MOTA	5845			11.104	79.213	
ATOM	5846		71.896	11.556	80.164	1.00 38.08
ATOM	5847		71.924	8.883	78.733	1.00 33.62
MOTA	5848	O PRO B 354	70.894	8.643	79.366	1.00 24.82
ATOM	5849	N TRP B 355	72.833	7.954	78.468	1.00 31.76
MOTA	5850		72.635	6.611	78.969	1.00 30.01
	5851	CB TRP B 355	73.653	5.655	78.359	1.00 34.02
MOTA		CG TRP B 355		4.378	77.910	1.00 44.37
ATOM	5852					
MOTA	5853	CD2 TRP B 355	73.263	3.072	78.436	1.00 45.39
MOTA	5854	CE2 TRP B 355	72.418	2.177	77.734	1.00 44.31
MOTA	5855	CE3 TRP B 355	74.107	2.569	79.432	1.00 47.19
ATOM	5856	CD1 TRP B 355	72.073	4.230	76.935	1.00 42.18
ATOM	5857	NE1 TRP B 355	71.704	2.910	76.826	1.00 37.84
ATOM	5858	CZ2 TRP B 355	72.395	0.808	77.999	1.00 44.97
	5859	CZ3 TRP B 355	. 74.084	1.207	79.694	1.00 50.83
MOTA				0.341	78.979	1.00 48.73
ATOM	5860		73.231			
MOTA	5861	C TRP B 355	72.819	6.685	80.485	1.00 30.87
ATOM	5862	O TRP B 355	73.622	7.474	80.981	1.00 26.93
ATOM	5863	N ARG B 356	72.061	5.880	81.218	1.00 24.96
ATOM	5864	CA ARG B 356	72.147	5.848	82.671	1.00 23.57
ATOM	5865	CB ARG B 356	70.811	6:319	83.257	1.00 24.71
	5866	CG ARG B 356	70.534	7.795	82.941	1.00 23.66
MOTA	5867	CD ARG B 356	69.067	8.212	83.055	1.00 20.14
ATOM						1.00 20.59
MOTA	5868	NE ARG B 356	68.926	9.610	82.642	
ATOM	5869	CZ ARG B 356	67.787	10.192	82.288	1.00 25.41
ATOM	5870	NH1 ARG B 356	66.644	9.508	82.287	1.00 17.01
MOTA	5871	NH2 ARG B 356	67.796	11.464	81.910	1.00 20.07
ATOM	5872	C ARG B 356	72.481	4.410	83.085	1.00 26.57
MOTA	5873	O ARG B 356	71.610	3.641	83.485	1.00 23.02
ATOM	5874	N GLY B 357	73.761	4.063	82.978	1.00 23.92
	5875	CA GLY B 357	74.186	2.712	83.294	1.00 25.54
ATOM	5876		74.796	2.464	84.657	1.00 24.35
ATOM						
ATOM	5877	O · GLY B 357	74.523.	3.161	85.628	1.00 25.88
ATCM	5878	N GLY B 358	75.638	1.444	84.718	1.00 24.32
ATOM	5879	CA GLY B 358	76.282	1.070	85.960	1.00 23.56
ATCM	5880	C GLY B 358	76.412	-0.441	85.924	1.00 29.26
ATOM	5881	O GLY B 358	76.146	-1.051	84.889	1.00 23.71
ATOM	5882	N GLU B 359	76.814	-1.051	87.033	1.00 27.64
	5883	CA GLU B 359		-2.503		
ATOM		CB GLU B 359	77.822	-2.936	88.265	1.00 30.40
ATOM	5884					
ATOM	5885	CG GLU B 359	77.125	-2.772	89.601	1.00 31.23
ATOM	5886	CD GLU B 359	77.844	-3.479	90.741	1.00 37.96
ATCM	5887	OE1 GLU B 359	77.287	-3.521	91.861	1.00 33.89
ATOM	5888	OE2 GLU B 359	78.959	~3.990	90.520	1.00 37.40
	5889	C GLU B 359	75.571	-3.122	87.261	1.00 31.35
ATOM	5890	O GLU B 359	74.612	-2.429	87.588	1.00 25.15
ATOM					87.053	1.00 29.61
ATOM	5891	N VAL B 360	75.482	-4.428		
ATCM	5892	CA VAL B 360	74.230	-5.147	87.251	1.00 26.21
ATOM	5893	CB VAL B 360	74.035	-6.270	86.200	1.00 28.47
ATOM	5894	CG1 VAL B 360	72.764	-7.045	86.492	1.00 22.74
ATOM	5895	CG2 VAL B 360	73.969	-5.670	84.796	1.00 29.70
	5896	C VAL B 360	74.342	-5.784	88.625	1.00 26.00
ATOM	5897	O VAL B 360	75.150	-6.693	88.821	1.00 27.55
ATCM	,00,	- 490 0 200		-0.023		

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Figure 18-90

ATOM	5898	N AR	G B	361	73	.553	-5.289	89.575	1.00	26.45
							-5.821	90.935		
MOTA	5899			361		.558				28.47
MOTA	5900			361	72	.479	-5.146	91.787		30.55
MOTA	5901	CG AR	G B	361	72	.937	-3.877	92.485	1.00	32.61
	5902		G B	361		.749	-3.163	93.117	1 00	35.00
ATOM										
ATOM	5903			361		.858	-2.617	92.094		30.31
ATOM	5904	CZ AR	GΒ	361	69	. 753	-1 <i>.</i> 925	92.350	1.00	29.45
MOTA	5905	NH1 AR	GB	361	69	.385	-1.689	93.605	1.00	18.49
						.041		91.348		30.49
ATOM	5906			361			-1.428			
MOTA	5907	C AR	G B	361	73	.351	-7.322	91.001		30.17
ATOM	5908	O AR	G B	361	72	.665	-7.910	90.168	1.00	23.60
ATOM	5909			362	73	.949	-7.922		. 1.00	33.09
							-9.351	92.272		36.94
MOTA	5910			362		.864				
MOTA	5911	CB LY	S B	362	74	. 687	-9.706	93.513	1.00	40.24
ATOM	5912	CG LY	SB	362	76	.190	-9.527	93.337	1.00	52.55
ATOM	5913		S B		76	.571	-8.126	92.849	1.00	56.65
								93.819		53.39
MOTA	5914			362		.149	-7.032			
MOTA	5915	NZ LY	S B	362	76	. 553	-5.680	93.341		48.87
ATOM	5916	C LY	5 B	362	72	.427	-9.826	92.463	1.00	32.84
ATOM	5917			362			-10.867	91.938	1.00	28.27
										34.67
ATOM	5918		JB	363		. 628	-9.075	93.215		
MOTA	5919	CA GL	JB	363	70.	. 245	-9.493	93.435		35.72
MOTA	5920	CB GL	JB	363	69.	.519	-8.532	94.390	1.00	36.04
ATOM	5921		JB	363		. 502	-7.077	93.977	1.00	44.81
			JB			859	-6.186	95.033		52.14
ATOM	5922			363						
MOTA	5923		JB	363		661	-6.370			48.46
MOTA	5924	OE2 GL	J B	363	69.	. 562	-5.300	95.566		57.31
MOTA	5925	C GLI	JB	363	69.	501	-9.619	92.111	1.00	30.68
	5926			363		695	-10.530	91.944		30.45
MOTA										26.19
MOTA	5927			364		784	-8.724	91.166		
MOTA	5928	CA VAI	ЪВ	364		138	-8.789	89.852		24.65
MOTA	5929	CB VAI	В	364	69.	536	-7.599	88.958	1.00	23.49
ATOM	5930	CG1 VAI	. R	364		924	-7.770	87.563	1.00	21.01
							-6:293	89.587		23.08
MOTA	5931	CG2 VAI		364		049				
ATOM .	5932	C VAI	ЪВ	364	69.	530	-10.083	89.144		23.19
ATOM	5933	O VAI	. B	364	68.	691	-10.749	88.542	1.00	23.06
ATOM	5934	N LYS	5 B	365	70.	810	-10.436	89.216	1.00	27.15
				365			-11.668	88.594		29.18
MOTA	5935									
ATOM	5936			365		821	-11.758	88.704		28.61
ATOM	5937	CG LYS	В	365	73.	554	-10.617	88.030	1.00	30.27
ATOM	5938	CD LYS	В	365	75.	074	-10.768	88.154	1.00	32.58
	5939			365		790	-9.587	87.516	1.00	29.13
MOTA								87.606		35.17
MOTA	5940			365		271	-9.689			
ATOM .	5941	C LYS	3 B	·365··	70.	666	-12.879	89.276		25.30
ATOM	5942	O LYS	В	365	70.	282	-13.837	88613	1.00	26.81
ATOM	5943	N ASE	В	366	70	559	-12.831	90.604	1.00	26.10
	-			366			-13.938	91.347		28.29
MOTA	5944									29.44
MOTA	5945	CB ASE		366		105	-13.731	92.859		
MOTA	5946	CG ASF	В	366	71.	557	-13.669	93.311		32.95
ATOM	5947	OD1 ASF	В	366	72.	446	-14.099	92.551	1.00	26.37
	5948	OD2 ASF					-13.216	94.442		35.26
MOTA								90.986		
MOTA	5949			366			-14.110	-		28.61
MOTA	5950	O ASP	В	366	68.	000	-15.231	90.869	1.00	27.00
ATOM	5951	N THR	В	367	67.	777	-13.002	90.801	1.00	28.63
							-13.080	90.438		27.35
MOTA	5952	CA THR		207						27.63
ATOM	5953	CB THR					-11.683	90.359		
ATOM	5954	OG1 THR	В	367	65.	771	-11.068	91.656		28.12
ATOM	5955	CG2 THR					-11.786	89.890	1.00	22.94
							-13.782	89.094		25.46
MOTA	5956									
ATOM	5957	O THR					-14.693	88.964		24.48
ATOM	5958	N LEU	В	368	66.	962	-13.361	88.092		23.33
ATOM	5959	CA LEU					-13.990	86.785	1.00	28.99
							-13.256	85.759		27.67
ATOM	5960									29.47
ATOM ·	5961	CG LEU					-12.070	85.046	1.00	47.4/
MOTA	5962	CD1 LEU			65.	923	-12.607	84.195	T.00	32.45
ATOM	5963	CD2 LEU	В	368	66.	546	-11.027	86.043	1.00	19.43
W 7 C/1.7				-	•	-		_		

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Figure 18-91

										•		
MOTA	5964	С	LE	U E	368		67.26	2 .	-15.454	86.888	1.00	32.40
ATOM	5965	0	L.E.	C E	3 3 6 8				-16.309			31.80
ATOM	5966	N	GL;	UE	3 3 6 9				-15.735			33.59
MOTA	5967	CA	GL:	J E	369				-17.101			39.68
ATOM	5968	CE	GL:	J E	3 3 6 9				-17.141			42.61
ATOM	5969				3 3 6 9				-17.138			50.65
ATOM	5970				369				-18.443			55.81
ATOM	5971				3 3 6 9				-18.561			57.42
ATOM	5972				369				-19.353			58.37
ATOM	5973				369				-17.954			34.94
ATOM	5974				369				-18.974			32.71
ATOM	5975				370				-17.541			34.92
ATOM	5976				370			_	-18.295			35.12
ATOM	5977	CB							-17.679			37.39
ATOM	5978	CG			370				-17.916			44.70
ATOM	5979	CD			370				-16.741			48.10
ATOM	5980	CE							-15.537			47.82
MOTA	5981	NZ							-14.387			45.41
ATOM	5982	C			370				-18.375			33.21
MOTA	5983	ō			370				-19.409			29.52
ATOM	5984	N			371		64.390		17.288			31.62
ATOM	5985	CA			371				17.274			37.19
ATOM	5986	CB	ALA			-			15.938			35.65
ATOM	5987	C			371	-	63.572		18.431			37.79
ATOM	5988	ō			371		62.627		19.137			34.45
ATOM	5989	Ŋ			372		64.810		18.644			40.10
ATOM	5990	CA			372				19.698			40.46
ATOM	5991	C			372		.64.746					43.15
ATOM	5992	ō			372				22.053	84.591		43.57
MOTA	5993	CB			372				19.694	84.517		40.51
ATOM	5994	CG			372				18.925	83.248		20.00
ATOM	5995	CD	LYS		372				19.390	82.635		20.00
ATOM	5996	CE	LYS		372		68.544		20.907	82.706		20.00
ATOM	5997	NZ	LYS		372 .				21.354	82.116		20.00
ATOM	5998	N	ALA		373		64.412		21.159	86.624		47.80
ATOM	5999	CA			373		64.014		22.425	87.239		49.71
ATOM	6000	CB	ALA	В	373		64.762		22.639	88.546		48.25
ATOM	6001	С	ALA	В	373		62.515	. –	22.443	87.494	1.00	53.38
MOTA	6002	0	ALA	В	373		61.844		23.313			58.01
ATOM	6003	OXI	' Ala	В	373		62.029		21.589	88.269	1.00	55.13
HETATM	2991	ZN	zn	С	1		49.660)	9.211	109.302	1.00	32.54
HETATM	2992	01	TSA	D	2		47.669	1	8.189	109.464	1.00	28.76
HETATM	2993	02	TSA	D	2		49.952		6.981	108.340	1.00	25.81
HETATM	2994	03	TSA	D	2		52.458		5.101	101.667	1.00	36.93
HETATM	2995	Nl	TSA		2		47.800		7.789	108.131	1.00	31.21
HETATM	2996	N2	TSA		2		53.013		-1.329	101.259	1.00	30.57
HETATM		C1	TSA		2		51.859		2.799	101.610	1.00	28.47
HETATM		C2	TSA		2		50.907		1.769	101.666		25.57
HETATM	2999	C3	TSA		2		51.241		0.419	101:551	1.00	21.68
HETATM		C4	TSA		2		52.626		0.026	101.366	1.00	23.11
HETATM		C5	TSA	D	2		53.589			101.303		25.02
HETATM	3002	C6	TSA		2		53.218		2.408	101.418		29.24
HETATM		C7	TSA		2		51.572			101.734		32.98
HETATM		C8	TSA	D	2		50.108		4.726	101.996	1.00	29.05
HETATM		C9	TSA	D	2		50.052		5.421	103.338		28.13
HETATM	3006	C10	TSA	D	2		49.060		5.357	104.279	1.00	25.99°
HETATM		C11	TSA		2		49.315		6.155	105.504		32.05
HETATM		C12	TSA		2		48.515			106.595		27.37
HETATM			TSA		2		48.855		6.994	107.756		29.02
HETATM	3010		TSA		2		49.680		5.693	100.864		30.21
HETATM	3011		TSA		2		47.776			104.132		30.60
HETATM	3012		TSA		2		54.438		-1.703	101.139		23.45
HETATM	3013	C16	TSA	D	2		52.044	-		101.316		23.15
HETATM	6004	ZN		E	1		52.949		1.842	85.681		28.19
HETATM	6005	01	TSA :		2		50.964		0.911	85.428		24.72
HETATM		02	TSA :	F	2		51.255		3.324	86.654	1.00	30.24

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HETAT	M 600	7 03	TS.	A F	2		5	1.569	9 6.5	12 02	.219	1 0		- 00
_	M 600			A F				3.34		11 00			27	
	M 600			A F							.634		0 27	
	M 601			a r				7.06			.713		16	
	M 601							.443			.304		27	
				A F				3.035	-	_	.267		25	
	M 6012			A F				7.227		57 93	.398		24	
	M 6013		TS				47	7.837	7 9.97	71 93	.583	1.00	25	.75
	M 6014		TS?	A F	2		49	.274	10.03	L7 93	.626		26	
HETAT	M 6015	C6	TSF	₹ F	2		50	.041	8.86	9 93	. 495	1.00	28	36
HETAT	M 6016	5 C7	TSA	A F	2		50	.349			.167	1.00		.27
HETAT	M 6017	7 C8	TSA	F	2			.716			.905		24	
HETAT	M 6018	C9	TSA	A F	2 -			.134			.518		27	
HETAT	M 6019				2			.419						
	M 6020		L TSA		2			.118			. 616	1.00		
	M 6021			_	2						.327	1.00		
	M 6022							.762			409	1.00		
					2			.529			170	1.00		
	M 6023				2			.208			994	1.00	28	. 83
	M 6024		TSA		2			.013		0 90.	863	1.00	26	.16
	M 6025				2		47	.699	12.45	6 93.	883	1.00	27	.37
	M 6026				2		45	.610	11.10	7 93.	679	1.00	25	.36
HETAT	M 6027	OH2	WAT	G	1		61	.391	6.72	3 88.	062	1.00		
HETAT	M 6028	OH2	TAW	G	2		55	.595	-4.44		558	1.00		. 53
HETATI	M 6029	OH2	WAT	G	3		58	.656		1 106.		1.00		
HETATI	4 6030	OH2	WAT	G	4			.347				1.00		
HETATI	4 6031	OH2			5			. 523	13.62	-	224	1.00		
	1 6032	OH2			6			. 466	-6.06		688			
	1 6033	OH2			7				-17.74			1.00	22.	. 41
	1 6034		WAT		8						769	1.00		
HETAT		OH2			9			.344			809	1.00		
HETATI			WAT						-14.90		717	1.00		
-					10			.540	-7.62			1.00		
HETATI			WAT-	_	11			414	-2.49		029	1.00		
HETAT		OH2			12			671		4 114.		1.00	32.	15
HETATM		OH2			13			335	10.67	9.117.	140	1.00	19.	47
HETATM			TAW	G	14		45.	565	9:46	9 79.	366	1.00	18.	81
HETATM	6041	OH2	WAT	G	15		43.	311	8.23	779.	508	1.00	26.	11
HETATM	6042	OH2	WAT	G	16		46.	628	13.883	3 104.		1.00		
HETATM	6043		WAT		17	•	40.	672	2.50			1.00		
HETATM	6044	OH2	WAT	G	18		61.	830	10.923			1.00		
HETATM	6045		WAT		19			813	0.831			1.00		
HETATM	6046	OH2	TAW	G	20			885	5.660			1.00		
HETATM	6047	OH2	WAT	G	21			382	-8.352				17.	
HETATM			WAT		22				-10.091					
HETATM			WAT		23			802	-3.446			1.00		
HETATM			WAT		24			292	12.112			1.00		
HETATM			WAT		25			747	8.830			1.00		
HETATM				G	26							1.00		
HETATM			WAT					952	9.79			1.00		
HETATM					27			268	2.80	106.		1.00		
				G	28			342	17.79:			1.00		
HETATM			WAT		29		72.		-6.985			1.00		
HETATM			WAT		30		39		9.257			1.00	22.	61
HETATM			WAT (31		61.	221	14.462	87.2	256	1.00	29.	85
HETATM		OH2	TAW	G	32		38.	167	22.692	107.4		1.00	36.4	40
HETATM		OH2	WAT (G	33		64.	657	-2.682	96.2		1.00		
HETATM	6060	CH2	WAT (3	34		44.	059	-2.698			1.00		
HETATM	6061	OH2	WAT (3	35		38.4		4.763			1.00		
HETATM	6062		WAT C		36		57.8		7.654			1.00		
HETATM	6063	OH2			37		57.0		3.145					
HETATM	6064	OH2			38		52.1		-1.400			1.00		
HETATM	6065	OH2			39							1.00		
							69.4		14.200	123.3		1.00		
HETATM		OH2 :			40		24.0		6.540	79.8		1.00		
HETATM	000/	OH2			41		46.6		-10.880	89.4		1.00		
HETATM	6068	OH2 V	VAT G		42		24.9		13.489			1.00	46.3	4
MTATE	6069	OH2 V	VAT G	•	43		46.5		-4.511	94.7		1.00	23.1	1
HETATM	6070	OH2 V	VAT G		14		51.4		13.833	86.3	06	1.00	27.0	18
HETATM	6071	OH2 W	VAT G	. 4	15		70.5	78	4.183	105.2	48	1.00		
HETATM.	6072	OH2 V	VAT G	; 4	16		53.9		-9.936			1.00		
									-	_				

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```
-0.443 63.035
                                                           1.00 28.35
                                  38.458
              OH2 WAT G
                          47
HETATM 6073
                                                           1.00 34.46
                                  64.786
                                           7.930 107.466
HETATM 6074
              OH2 WAT G
                          48
                                          36.521 114.809
                                                            1.00 40.51
                                  50.823
                          49
              OH2 WAT G
HETATM 6075
                                                            1.00 39.11
                                                   68.080
                          50
                                  33.963 -10.352
              OH2 WAT G
HETATM 6076
                                                            1.00 33.30
                                  71.328 -14.321
                                                   86.007
HETATM 6077
              OH2 WAT G
                          51
                                  63.272 10.210
                                                            1.00 35.75
                                                   79.836
                          52
              OH2
                  WAT G.
HETATM 6078
                                                            1.00 29.57
                                                   94.306
                                  59.263 -12.096
              OH2 WAT G
                          53
HETATM 6079
                                                            1.00 27.97
                                  46.041 10.641
                                                   76.561
              OH2 WAT G
                          54
HETATM 6080
                                  46.614 -13.620
                                                   89.775
                                                            1.00 24.25
                          55
              OH2 WAT G
НЕТАТМ 6081
                                                   89.097
                                                            1.00 29.19
                                  76.600
                                            0.622
                          56
              OH2_WAT G
HETATM 6082
                                                            1.00 34.05
                                            6.439
                                                   79.089
                          57
                                  53.555
              OH2 WAT G
HETATM 6083
                                                   83.310
                                                            1.00 35.02
                                  71.301
                                           11.026
              OH2 WAT G
                          58
HETATM 6084
                                                   81.594
                                                            1.00 33.21
                                  28.188
                                           -9.956
                          59
              OH2 WAT G
HETATM 6085
                                                            1.00 27.64
                                           20.992
                                                   98.483
                                  53.084
              OH2
                  WAT G
                          60
HETATM 6086
                                                   93.423
                                                            1.00 30.30
                                            8.630
                                  59.484
              OH2 WAT G
                          61
HETATM 6087
                                           -3.809
                                                   95.805
                                                            1.00 33.04
                                  26.195
              OH2 WAT G
                          62
HETATM 5088
                                                   89.620
                                                            1.00 37.39
                                  26.095
                                           -0.121
              OH2 WAT G
                          63
HETATM 6089
                                           -6.141 109.711
                                                            1.00 20.88
                                  47.100
HETATM 6090
              OH2 WAT G
                          64
                                                            1.00 30.38
                                            0.731
                                                  92.275
              OH2
                  WAT G
                          65
                                  23.273
HETATM 6091
                                                            1.00 37.51
                                                   72.694
                                  45.340 -24.751
HETATM 6092
              OH2 WAT G
                          66
                                          16.234 111.676
                                                            1.00 34.63
                          67
                                  33.754
HETATM 6093
              OH2 WAT G
                                                            1.00 47.11
                                           19.209 126.276
                  WAT G
                          68
                                  52.831
              OH2
HETATM 6094
                                           16.953 111.099
                                                            1.00 26.24
                          69
                                  50.218
                  WAT G
HETATM 6095
              OH2
                                                            1.00 24.95
                                  44.791
                                            5.844
                                                   70.857
              OH2 WAT G
                          70
HETATM 6096
                                                   82.921
                                                            1.00 29.48
                                  49.517 -18.731
                          71
HETATM 6097
                  WAT G
              OH2
                                          10.131 116.550
                                                            1.00 48.70
                          72
                                  76.379
              OH2
                  WAT G
HETATM 6098
                                                   87.873
                                                            1.00 46.35
                          73
                                  30.214
                                           -8.086
              OH2 WAT G
HETATM 6099
                                                            1.00 30.80
                          74
                                   45.320
                                           12.061
                                                   80.458
              OH2 WAT G
HETATM 6100
                                  72.881
                                            5.360
                                                   86.249
                                                            1.00 29.04
HETATM 6101
              OH2 WAT G
                          75
                                  59.674 -23.046
                                                            1.00 41.96
                                                   87.252
                          76
              OH2
                  WAT G
HETATM 6102
                                                            1.00 26.45
                                            7.921 100.345
                          77
                                   40.619
                  WAT G
HETATM 6103
              OH2
                                                            1.00 36.27
                                   41.666 -19.477
                                                   70.073
                          78
HETATM 6104
              OH2 WAT G
                                  46.408
                                                   92.717
                                                            1.00 25.78
                                          -6.539
                  WAT G
                          79
HETATM 6105
              OH2
                                                            1.00 28.34
                                                   81.646
                                  35.743 -12:230
              OH2 WAT G
                          80
HETATM 6106
                                                            1.00 41.15
                                            8.745 121.961
                          81
                                  28.268
              OH2 WAT G
HETATM 6107
                                                            1.00 32.34
                                  68.843
                                            3.154
                                                   71,986
HETATM 6108
              OH2
                  WAT G
                          82
                                   52.125 -11.158
                                                    85.150
                                                            1.00 24.14
              OH2
                  WAT G
                          83
HETATM 6109
                                                   92.264
                                                            1.00 26.12
                                           -1.773
                                  75.374
              OH2 WAT G
                          84
HETATM 6110
                                                            1.00 37.07
                                   46.957
                                           12.230 142.271
                          85
HETATM 6111
              OH2
                  WAT G
                                                            1.00 55.58
                                  63.789
                                            9.551
                                                    64.329
HETATM 6112
              OH2
                  WAT G
                          86
                                                            1.00 58.55
                                           21.185
                                                    72.215
                                   60.672
                          87
              OH2
                  WAT G
HETATM 6113
                                                    82.064
                                            9.505
                                                            1.00 31.10
                          88
                                   56.547
                  WAT G
 нетатм 6114
              OH2
                                                            1.00 29.70
                                                    92.250
                                  26.366
                                           -0.876
                          89
HETATM 6115
              OH2 WAT G
                                                            1.00 32.85
                                          -16.583
                                                    80.808
                                   67.504
                  WAT G
                          90
 HETATM 6116
              OH2
                                            1.899
                                                            1.00 42.95
                                                    82.068
                          91
                                  23.910
              OH2
                  WAT G
 HETATM 6117
                                            4.106 117.380
                                                            1.00 30.05
                                   50.032
              OH2 WAT G
                          92
HET/ TM 6118
                                                            1.00 43.59
                                           -9.492
                                                   83.952
                                  26.774
 HET. IM 6119
              OH2
                  WAT G
                          93
                                                  113.787
                                                            1.00 40.17
                                           -0.637
                                   42.714
                          94
              OH2
                  WAT G
 HETA.M 6120
                                            7.989 134.170
                                                            1.00 47.82
                          95
                                  57.966
              OH2 WAT G
 HETATM 6121
                                                            1.00 36.62
                                           -3.550 119.086
                                   54.478
                          96
 HETATM 6122
              OH2 WAT G
                                   53.065
                                           11.696 101.718
                                                            1.00 41.62
                  WAT G
                          97
              OH2
 HETATM 6123
                                                    68.207
                                                            1.00 45.98
                                          -23.645
                                   58.286
                  WAT G
                          98
 HETATM 6124
              OH2
                                           -9.614 121.975
                                                            1.00 34.57
                                   54.855
                          99
              OH2
                  WAT G
 HETATM 6125
                                                    57.145
                                                            1.00 42.14
                                   57.408
                                           -3.352
                  WAT G 100
              OH2
 HETATM 6126
                                           20.353 123.667
                                                            1.00 33.87
                                   63.590
                  WAT G 101
 HETATM 6127
              OH2
                                                    72.392
                                                            1.00 30.23
                                          -23.143
                                   48.129
                  WAT G 102
              OH2
 HETATM 6128
                                                            1.00 52.01
                                                    76.094
                                   62.834
                                            6.913
              OH2
                  WAT G 103
 HETATM 6129
                                            6.529
                                                    73.089
                                                            1.00 36.29
                                   34.566
                  WAT G 104
              OH2
 HETATM 6130
                                                    67.459
                                                            1.00 36.85
                                   51.588
                  WAT G 105
                                           20.869
              OH2
 HETATM 6131
                                                            1.00 42.87
                                                   129.379
                                           18.020
                                   28.160
                  WAT G 106
              OH2
 HETATM 6132
                                          -11.452
                                                    57.603
                                                            1.00 43.62
                  WAT G 107
                                   49.082
 HETATM 6133
              OH2
                                                    93.281
                                                            1.00 41.95
                  WAT G 108
                                   44.717
                                           -8.605
              OH2
 HETATM 6134
                                                            1.00 35.71
                                          -11.900
                                                    94.019
                                   67.088
              OH2
                  WAT G 109
 HETATM 6135
                                                            1.00 35.40
                                           22.763 100.800
                  WAT G 110
                                   49.561
              OH2
 HETATM 6136
                                                            1.00 56.20
                                           10.960 124.536
                                   75.853
                  WAT G 111
 HETATM 6137
              OH2
                                                            1.00 36.40
                                   54.383
                                            8,930 136.095
              OH2 WAT G 112
 HETATM 6138
```

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:

```
33.114
42.618
 HETATM 6139
               OH2 WAT G 113
                                            1.764 67.443
                                                            1.00 37.01
 HETATM 6140
                                           -4.357 102.345
               OH2 WAT G 114
                                                             1.00 39.18
               OH2 WAT G 115
 HETATM 6141
                                   53.605
                                          -10.816
                                                   66.281
                                                            1.00 31.62
 HETATM 6142
               OH2
                   WAT G 116
                                   73.410
                                            -1.010
                                                   90.400
                                                            1.00 34.72
               OH2 WAT G 117
                                   68.899
 HETATM 6143
                                             3.789 110.221
                                                            1.00 35.69
 HETATM 6144
               OH2 WAT G 118
                                   31.474
                                           19.159 112.425
                                                            1.00 28.08
               OH2 WAT G 119
                                   39.749
 HETATM 6145
                                            -0.616 132.457
                                                            1.00 37.43
 HETATM 6146
               OH2 WAT G 120
                                   44.921
                                             1.089 137.137
                                                            1.00 40.80
               OH2 WAT G 121
                                   31.081
 HETATM 6147
                                             7.617
                                                   75.105
                                                            1.00 40.86
 HETATM 6148
                   WAT G 122
               OH2
                                   35.554
                                           12.017 105.965
                                                            1.00 33.58
 HETATM 6149
               OH2 WAT G 123
                                   41.381
                                                    70.872
                                           -23.534
                                                            1.00 38.10
 HETATM 6150
               OH2
                   WAT G 124
                                   31.999
                                            1.992
                                                    73.813
                                                            1.00 33.97
 HETATM 6151
               OH2 WAT G 125
                                   55.761
                                                            1.00 47.66
                                           10.285 101.654
 HETATM 6152
               OH2 WAT G 126
                                   30.596
                                           12.964 133.642
                                                            1.00 37.98
 HETATM 6153
               OH2 WAT G 127
                                   59.611
                                            5.347 136.114
                                                            1.00 46.39
 HETATM 6154
               OH2 WAT G 128
                                   24.190
                                           12.220 124.679
                                                            1.00 30.77
 HETATM 6155
               OH2 WAT G 129
                                   70.078
                                            4.455
                                                    86.283
                                                            1.00 36.11
 HETATM 6156
               OH2 WAT G 130
                                   57.882
                                           -4.314 125.597
                                                            1.00 41.40
               OH2 WAT G 131
 HETATM 6157
                                   45.838 -20.690
                                                    65.884
                                                            1.00 35.98
 HETATM 6158
               OH2 WAT G 132
                                   47.574
                                            3.186
                                                    79.027
                                                            1.00 36.67
 HETATM 6159
               OH2 WAT G 133
                                   46.856
                                          -18.901
                                                    62.295
                                                            1.00 45.40
 HETATM 6160
               OH2 WAT G 134
                                   40.164
                                            5.047
                                                    95.358
                                                            1.00 31.38
               OH2 WAT G 135
 HETATM 6161
                                   27.268
                                           -0.405
                                                  122.461
                                                            1.00 38.16
 HETATM 6162
              OH2 WAT G 136
                                   54.200 -20.155
                                                    66.212
                                                            1.00 37.55
 HETATM 6163
              OH2 WAT G 137
                                   45.435 -10.534
                                                            1.00 37.96
                                                  103.626
 HETATM 6164
              OH2 WAT G 138
                                   31.633
                                           25.030 106.499
                                                            1.00 43.94
              OH2 WAT G 139
 HETATM 6165
                                   79.029
                                           -7.518
                                                   93.606
                                                            1.00 40.55
              OH2 WAT G 140
 HETATM 6166
                                  68.597
                                           20.711 111.685
                                                            1.00 33.25
 HETATM 6167
              OH2 WAT G 141
                                  64.263
                                            8.524 113.832
                                                            1.00 40.63
 HETATM 6168
              OH2 WAT G 143
                                  49.387 -24.485
                                                   70.152
                                                            1.00 34.07
 HETATM 6169
              OH2 WAT G 144
                                  23.383
                                           -3.854
                                                    83.604
                                                            1.00 32.22
 HETATM 6170
              OH2 WAT G 145
                                  42.360
                                           -0.710
                                                    61.686
                                                            1.00 35.94
              OH2 WAT G 146
 HETATM 6171
                                                    65.685
                                  34.421
                                           -3.304
                                                            1.00 35.42
HETATM 6172
              OH2 WAT G 147
                                  31.506
                                            3.409
                                                    89.579
                                                            1.00 39.86
HETATM 6173
              OH2 WAT G 148
                                  34.963
                                                            1.00 31.12
                                           10.688
                                                   91.806
              OH2 WAT G 149
HETATM 6174
                                  54.859 -15.085
                                                   96.769
                                                            1.00 46.65
HETATM 6175
              OH2 WAT G 150
                                  34.695
                                            2.391 131.273
                                                            1.00 39.22
HETATM 6176
              OH2 WAT G 151
                                  40.348
                                            1.395
                                                  61.905
                                                            1.00 34.09
HETATM 6177
                                  66.912
                                           17.666 127.489
              OH2 WAT G 152
                                                            1.00 45.19
HETATM 6178
              OH2 WAT G 153
                                  31.096
                                          19.900 103.232
                                                            1.00 43.45
HETATM 6179
              OH2 WAT G 154
                                                            1.00 28.86
                                  28.074
                                           -4.222
                                                   70.175
HETATM 6180
              OH2 WAT G 155
                                  63.586
                                          -1.894
                                                   99.003
                                                            1.00 41.15
                                  54.145 -22.222
HETATM 6181
              OH2 WAT G 156
                                                   88.415
                                                            1.00 40.92
HETATM 6182
              OH2 WAT G 157
                                  62.443
                                           13.765
                                                   89.547
                                                            1.00 33.69
HETATM 6183
              OH2 WAT G 158
                                           9.798 101.311
                                  58.832
                                                            1.00 31.00
                                          -5.528 119.322
HETATM 6184
              OH2 WAT G 159
                                  37.701
                                                            1 00 45.00
HETATM 6185
              OH2 WAT G 160
                                  43.599
                                          13.442 131.274
                                                            1 00 38.43
HETATM 6186
              OH2 WAT G 161
                                  23.540
                                          -1.137
                                                   96.111
                                                            1 00 51.83
HETATM 6187
              OH2 WAT G 162
                                  59.915
                                          -4.318 110.873
                                                           1.00 41.92
HETATM 6188
              OH2 WAT G 163
                                          -8.264
                                  51.265
                                                   60.546
                                                           1.00 31.25
HETATM 6189
             OH2
                 WAT G 164
                                  58.109
                                           7.024
                                                   98.294
                                                           1.00 46.30
HETATM 6190
             OH2
                 WAT G 165
                                          18.195
                                                           1.00 37.53
                                  46.553
                                                   74.179
             OH2 WAT G 166
HETATM 6191
                                  55.706 -21.025
                                                   92.515
                                                           1.00 43.91
HETATM 6192
             OH2 WAT G 167
                                  67.146
                                          -1.958 109.704
                                                           1.00 43.13
HETATM 6193
             OH2 WAT G 168
                                  47.445
                                          -3.047 134.746
                                                           1.00 27.99
HETATM 6194
             OH2 WAT G 169
                                           5.304
                                                   63.562
                                  65.193
                                                           1.00 36.05
HETATM 6195
             OH2 WAT G 170
                                  36.176
                                           8.979
                                                 102.024
                                                           1.00 39.63
                                           5.797
HETATM 6196
             OH2 WAT G 171
                                  70.527
                                                   70.886
                                                           1.00 44.69
HETATM 6197
                                  67.166
                                           8.735
             OH2 WAT G 172
                                                   74.628
                                                           1.00 51.41
HETATM 6198
             OH2 WAT G 173
                                  19.700
                                           9.630
                                                   81.850
                                                           1.00 53.49
                                 55.875
HETATM 6199
             OH2 WAT G 174
                                          11.277
                                                   87.176
                                                           1.00 38.63
HETATM 6200
             OH2 WAT G 175
                                 61.874
                                                   91.682
                                           8.432
                                                           1.00 40.08
HETATM 6201
             OH2
                 WAT G 176
                                 36.771
                                          -6.815 121.530
                                                           1.00 32.57
HETATM 6202
             OH2 WAT G 177
                                 63.224
                                           7.776
                                                 89.317
                                                           1.00 29.83
HETATM 6203
             OH2 WAT G 178
                                          15.345 132.470
                                 29.606
                                                           1.00 47.28
HETATM 6204
             OH2 WAT G 179
                                          11.799
                                                  98.957
                                 52.811
                                                           1.00 36.09
```

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```
1.00 36.19
                                  38.589
                                         18,249
                                                   88.356
              OH2 WAT G 180
HETATM 6205
                                                   61.135
                                                            1.00 34.24
                                  43.734 -15.681
              OH2 WAT G 181
HETATM 6206
                                                            1.00 37.96
                                  42.283
                                          15.251
                                                   91.437
HETATM 6207
                  WAT G 182
              OH2
                                                            1.00 45.78
                                  57.121 -11.129 126.206
              OH2
                  WAT G
                        183
HETATM 6208
                                                            1.00 36.55
                                  50.011 -19.367
                                                   92.127
                  WAT G 184
HETATM 6209
              OH2
                                  56.880
                                                   .95.969
                                                            1.00 39.12
                  WAT G 185
                                            2.453
HETATM 6210
              OH2
                                           14.125 125.052
                                                            1.00 32.68
                  WAT G 186
                                  26.356
              OH2
HETATM 6211
                                           20.230 122.650
                                                            1.00 45.67
                                  24.631
                  WAT G 187
HETATM 6212
              OH2
                                  23.516
                                                   81.599
                                                            1.00 42.16
                                            4.964
                  WAT G.188
HETATM 6213
              OH2
                                           14.964
                                                   62.948
                                                            1.00 50.18
                                  55.017
                  WAT G 189
              OH2
HETATM 6214
                                                            1.00 37.04
                                  33.371
                                           13.710 105.640
                  WAT G 190
HETATM 6215
              OH2
                                  44.466
                                                   91.144
                                                            1.00 36.62
                                          -10.386
                  WAT G 191
HETATM 6216
              OH2
                                           22.668 121.285
                                                            1.00 38.19
                                  28.437
                        192
              OH2
                  WAT G
HETATM 6217
                                                            1.00 42.05
                                           24.957 122.112
                                  29.786
                  WAT G 193
HETATM 6218
              OH2
                                            3.461
                                                            1.00 48.35
                                                   96.101
                                  28.852
                  WAT G 194
HETATM 6219
              OH2
                                           11.318
                                                   92.011
                                                            1.00 32.60
                  WAT G 195
                                   41.681
              OH2
HETATM 6220
                                                            1.00 47.70
                                   26.812 -10.229 111.631
HETATM 6221
                  WAT G 196
              OH2
                                                            1.00 48.86
              OH2 WAT G 197
                                   42.432 -23.250
                                                   76.629
HETATM 6222
                                          12.756 121.410
                                                            1.00 43.09
                                  25.484
                  WAT G 198
HETATM 6223
              OH2
                                   43.514 -20.514 111.706
                                                            1.00 46.80
                  WAT G 199
              OH2
HETATM 5224
                                                            1.00 44.89
                  WAT G 200
                                   74.273 -13.079
                                                   95.699
              OH2
HETATM 6225
                                   59.982
                                          24.381 103.984
                                                            1.00 40.63
HETATM 6226
                  WAT G 201
              OH2
                                   67.164 -12.771
                                                             1.00 35.13
                                                    74.705
                  WAT G 202
              OH2
HETATM 6227
                                                    79.238
                                                             1.00 33.53
                  WAT G 203
                                   28.708
                                            9.211
HETATM 6228
              OH2
                                                             1.00 48.49
                                   53.256
                                           -3.576 122.243
                        204
              OH2
                  WAT G
HETATM 6229
                                   50.706
                                                    87.357
                                                             1.00 41.26
                                           16.208
HETATM 6230
              OH2
                  WAT G 205
                                                    98.339
                                                             1.00 39.15
                                           34.998
                  WAT G
                        206
                                   50.000
              OH2
HETATM 6231
                                                             1.00 29.70
                  WAT G 207
                                   68.078
                                          -16.236
                                                    83,621
HETATM 6232
              OH2
                                                             1.00 50.82
                                   24.395
                                                   111.635
                                           -4.134
                  WAT G 208
HETATM 6233
              OH2
                                           -2.664 114.289
                                                             1.00 44.49
                                   53.384
              OH2
                  WAT G 209
HETATM 6234
                                                   94.788
                                                             1.00 31.97
                                   60.120
                                           -9.482
                  WAT G 210
HETATM 6235
              OH2
                                                             1.00 44.97
                                           17.472
                                                   111.744
                                   23.405
HETATM 6236
              OH2
                  WAT G 211
                                   46.214
                                                             1.00 59.14
                                           20.943
                                                    76.878
нетатм 6237
              OH2
                  WAT G 212
                                                    97.109
                                                             1.00 41.78
                                   29.754
                                            6:983
                  WAT G 213
              OH2
HETATM 6238
                                                             1.00 41.02
                                                    55.181
                                   46.820
                                           -0.465
              OH2 WAT G 214
HETATM 6239
                                                   124.775
                                                             1.00 38.42
                                           22.096
                                   59.143
нетатм 6240
              OH2
                  WAT G 215
                                   42.674
                                                    66.037
                                                             1.00 32.50
                                           14.088
                  WAT G 216
              OH2
HETATM 6241
                                                             1.00 56.50
                                                    98.186
                                           -7.248
                        217
                                   55.009
                  WAT G
              OH2
HETATM 6242
                                                   109.653
                                                             1.00 49.66
                                   63.361
                                           -8.209
                  WAT G 218
HETATM 6243
              OH2
                                                             1.00 50.91
                                           -8.146
                                                    94.671
                                   66.583
HETATM 6244
              OH2
                  WAT G 219
                                                             1.00 36.99
                                   44.627
                                           -2.583
                                                    93.919
                  WAT G 220
              OH2
HETATM 6245
                                                             1.00 47.24
                                           -8.606
                                                    79.502
                                   24.470
                  WAT G 221
НЕТАТМ 6246
              OH2
                                                    83.973
                                                             1.00 50.43
                                   76.913
                                           -7.777
                  WAT G 222
              OH2
HETATM 6247
                                                             1.00 42.47
                                                   129.136
                                   32.788
                                            0.651
                  WAT G 223
 HETATM 6248
              OH2
                                          -16.880
                                                    88.817
                                                             1.00 46.69
                                   73.731
                        224
                  WAT G
 HETATM 6249
              OH2
                                                    93.970
                                                             1.00 43.17
                                   78.567
                                           -2.802
                  WAT G 225
HETATM 6250
              OHE
                                                             1.00 35.84
                                                    57.532
                                            1.248
                                   45.681
                  WAT G 226
HETATM 6251
              OH.
                                           15.236
                                   38.263
                                                    84.711
                                                             1.00 42.39
                  WAT G 227
              OHL
 HETATM 6252
                                                   108.488
                                                             1.00 52.23
                  WAT G 228
                                   38.933
                                           35.224
HETATM 6253
              OH2
                                                             1.00 46.56
                                                    70.228
                                   33.755
                                           14.939
 нетатм 6254
              OH2
                  WAT G
                        229
                                           34.184
                                                   100.859
                                                             1.00 52.96
                                   51.521
                  WAT G 230
              OH2
 HETATM 6255
                                            0.565
                                                    63.039
                                                             1.00 31.02
                                   34.140
                  WAT G 231
HETATM 6256
              OH2
                                                    81.662
                                                             1.00 39.83
                                   37.277
                                           13.977
                        232
 HETATM 6257
              OH2
                  WAT G
                                                             1.00 28.55
                                            5.947
                                                    93.216
                                   57.307
                  WAT G 233
              OH2
 HETATM 6258
                                                             1.00 53.16
                                                   125.707
                                   31.718
                                           16.820
                  WAT G
                        234
HETATM 6259
              OH2
                                                   110.067
                                                             1.00 51.02
                                   60.624
                                           31.119
                  WAT G
                        235
              OH2
 HETATM 6260
                                                             1.00 57.50
                                   44.357
                                             4.267
                                                    94.916
                  WAT G 236
              OH2
 HETATM 6261
                                             1.072
                                                    70.111
                                                             1.00 38.65
                                   68.454
                  WAT G
                         237
 HETATM 6262
              OH2
                                                    79.253
                                                             1.00 44.95
                                   27.836
                                             6.773
                  WAT G
                        238
              OH2
 HETATM 6263
                                                             1.00 42.95
                                                    81.087
                                   54.933
                                           23.344
                  WAT G 239
              OH2
 HETATM 6264
                                                     71.579
                                                             1.00 45.78
                                           -15.271
                                   34.072
              OH2
                  WAT G
                         240
 HETATM 6265
                                                    61.973
                                                             1.00 38.91
                                   35.966
                                            -1.059
                         241
              OH2
                  WAT G
 HETATM 6266
                                                             1.00 44.85
                                                   127.376
                                   29.687
                                             1.898
                  WAT G
                        242
              OH2
 HETATM 6267
                                                   113.501
                                                             1.00 38.32
                                          -10.150
              OH2
                  WAT G 243
                                   49.534
 HETATM 6268
                                                             1.00 48.83
                                                     96.696
                                             9.773
                                   57.252
                  WAT G 244
 HETATM 6269
              OH2
                                                             1.00 38.54
                                                     80.972
                                   62.310
                                           13,262
              OH2 WAT G 245
 HETATM 6270
```

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HETAT	4 6271	OH2 WAT	' G 246	50.248	-5.552	102.815	1.00 43.23
HETATM	1 6272	OH2 WAT	G 247	47.966	21.564	79.321	1.00 36.79
HETATM		OH2 WAT	G 248	62.507	20.866	108.414	1.00 35.30
HETATM			G 249		19.763		1.00 47.49
HETATM	1 62/5		G 250		9.828		1.00 33.71
HETATM	6276	OH2 WAT	G 251	35.304	-6.179	66.319	1.00 36.23
HETATM		OH2 WAT	G 252	39.218	-12.667	85.010	1.00 36.17
HETATM			G 253		5.089		1.00 46.38
HETATM	1 02/3		G 254			122.119	1.00 55.07
HETATM	6280		G 255			128.600	1.00 30.64
HETATM	6281	OH2 WAT	G 256	- 32.019	-12.973	113.965	1.00 34.48
HETATM		OH2 WAT	G 257	54.081	3.421	56.994	1.00 39.11
HETATM	6283		G 258	32.801	-6.170		1.00 35.72
		OH2 WAT			0.301		1.00 36.57
HETATM				45.040			
HETATM			G 260	39.815		128.855	1.00 40.10
HETATM	6286	OH2 WAT	G 261	28.763	10.408	93.790	1.00 44.39
HETATM	6287	OH2 WAT	G 262	49.668	-12.050	60.539	1.00 50.89
HETATM	6288	OH2 WAT	G 263	64.353		117.495	1.00 62.67
HETATM	6280		G 264	75.183	13 021	128.124	1.00 50.42
HETATM	6205						
HETATM			G 265	46.289	6.826	52.485	1.00 46.86
HETATM	6291	OH2 WAT	G 266	68.708	13.973	70.958	1.00 37.90
HETATM	6292	OH2 WAT	G 267	71.504	12.997	130.029	1.00 38.78
HETATM	6293		G 268	36.309		130.364	1.00 42.92
UPTWIN	6204		G 269		12.195	79.625	1.00 42.92
HETATM	0294						
HETATM		OH2 WAT		71.952	13.021	74.292	1.00 37.70
HETATM	6296	OH2 WAT	G 271.	44.433	-17.578	62.734	1.00 49.33
HETATM		OH2 WAT	G 272	26.917	15.038	89,067	1.00 38.07
HETATM	6208		G 273	63.380		126.550	1.00 41.73
HETAIM	6290						
HETATM	6299	OH2 .WAT		63.360	-5.356	95.641	1.00 37.54
HETATM	6300	OH2 WAT		65.947	-13.015	97.485	1.00 37.42
HETATM	6301	OH2 WAT	G 276	26.406	25.831	117.328	1.00 48.37
HETATM		OH2 WAT	G 277	41.893	-10.251	98.201	1.00 46.36
HETATM	6303	OH2 WAT		30.343		117.764	1.00 49.87
HETAIM	6363						
HETATM		OH2 WAT			32:419		1.00 43.93
HETATM		OH2 WAT	G 280	50.553	-1.365	120.511	1.00 54.02
HETATM	6306	OH2 WAT	G 281	60.428	13.652	105.130	1.00 31.10
HETATM	6307	OH2 WAT	G 282	30.342	2.204	70.246	1.00 45.19
HETATM	6308	OH2 WAT		60.358	15.921		1.00 33.17
HETAIM	6300						
HETATM			G 284	64.193	3.421	62.117	1.00 45.81
HETATM	6310	OH2 WAT	G 285	45.468	6.113		1.00 48.98
HETATM	6311	OH2 WAT	G 286	47.514	3.808	98.279	1.00 46.45
HETATM		OH2 WAT	G 287	72.144	-6.345	77.930	1.00 40.04
HETATM		OH2 WAT		54.142	-5.100	99.674	1.00 43.62
REIAIM	6311			-	13.297	65.822	1.00 46.98
MTATM			G 289	48.982			
HETATM	6315		G 290	41.171	34.107		1.00 51.76
HETATM	6316	OH2 WAT	G 291	36.494	37.195	104.170	1.00 44.27
HETATM	6317	OH2 WAT	G 292	48.580	23.117	85.456	1.00 40.96
HETATM	6318		G 293	55.853	22.934	98.099	1.00 40.95
TELVIL	6310				11.077	89.427	1.00 41.21
HETATM	0313		G 294	61.720			
HETATM	6320	OH2 WAT				114.112	1.00 42.32
MTATH	6321	OH2 WAT	G 296	53.001	-6.305	129.052	1.00 37.41
HETATM		OH2 WAT	G 297	70.258	24.928	82.843	1.00 48.09
HETATM		OH2 WAT		77.493	0.940	130.507	1.00 51.77
RETAIM	6323						
HETATM	6524	OH2 WAT		32.233	12.182	83.028	1.00 53.51
HETATM	6325	OH2 WAT	G 300	40666	12.878	65.747	1.00 46.49
HETATM	6326	OH2 WAT	G 301	50.977	12.831	114.597	1.00 48.51
HETATM	6327	OH2 WAT		54.236	3.817	92.196	1.00 41.15
**********	6320	CH2 WAT		59.527	-1.343	107.471	1.00 36.71
HETATM	0320						
HETATM	6329	OH2 WAT		70.331	3.940	89.312	1.00 47.70
HETATM	6330	CH2 WAT	G 305	60.626	6.969	127.780	1.00 41.96
MTATH	6331	OH2 WAT	306	42.156	-0.139	133.156	1.00 32.19
HETATM	6332	OH2 WAT		58.886	16.514	99.413	1.00 53.60
ELATM	2222					96,570	1.00 40.36
HETATM	222	OH2 WAT		67.617	-1.589		
HETATM	6334 .	OH2 WAT			-10.936	98.849	1.00 48.80
HETATM	6335	OH2 WAT (310	45.576	25.388	131.914	1.00 48.99
METATM	6336	CH2 WAT		37.583	-6.243	64.257	1.00 37.06
				2,.000	-		

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Figure 18-97

```
16.408 94.600 1.00 45.07
                                  66.759
HETATM 6337
              OH2 WAT G 312
                                                           1.00 52.23
              OH2 WAT G 313
                                24.142
                                          11.212 113.340
HETATM 6338
                                                           1.00 39.88
                                  69.409
                                          16.702
                                                  64.230
HETATM 6339
              OH2 WAT G 314
                                                           1.00 50.23
                                          24.858 115.328
                                  22.064
              OH2 WAT G 315
HETATM 6340
                                           9.551 100.345
                                                           1.00 37.32
              OH2 WAT G 316
                                  50.171
HETATM 6341
                                                           1.00 44.78
                                          31.302 119.497
              OH2 WAT G 317
                                  55.104
HETATM 6342
                                                           1.00 44.21
                                  65.333 -10.105
              OH2 WAT G 318
                                                   95.866
HETATM 6343
                                  31.415
37.423
                                          -2.472
                                                 128.127
                                                           1.00 41.95
              OH2 WAT G 319
HETATM 6344
                                                           1.00 44.79
                                          13.143
                                                   88.069
              OH2 WAT G 320
HETATM 6345
                                                           1.00 54.69
                                                   96.509
                                          14.292
              OH2_WAT G 321
                                  43.619
HETATM 6346
                                          14.555
                                                 126.016
                                                           1.00 42.75
              OH2 WAT G 322
                                  68.048
HETATM 6347
                                  34.778
                                          -2.509 130.204
                                                           1.00 37.06
              OH2 WAT G 323
HETATM 6348
                                                           1.00 47.34
                                          18.144 103.841
              OH2 WAT G 324
                                  27.972
HETATM 6349
                                                           1.00 38.03
                                                  97.592
                                  53.550
                                          23.610
              OH2 WAT G 325
HETATM 6350
                                  33.776
                                           4.171 103.451
                                                           1.00 50.60
              OH2 WAT G 326
HETATM 6351
                                          35.632 114.870
                                                           1.00 48.34
              OH2 WAT G 327
                                  37.862
нетатм 6352
                                                           1.00 38.77
              OH2 WAT G 328
                                  50.893
                                          14.612
                                                   93.478
HETATM 6353
                                  71.422 -20.913
                                                   86.137
                                                           1.00 47.69
нетатм 6354
              OH2 WAT G 329
                                  50.310 -23.133
                                                   74.502
                                                           1.00 41.94
              OH2 WAT G 330
HETATM 6355
                                                           1.00 54.93
                                                   60.583
              OH2 WAT G 331
                                  41.520
                                           7.269
HETATM 6356
                                          13.737 106.089
                                  75.879
                                                            1.00 44.65
              OH2 WAT G 332
HETATM 6357
                                           9.027 138.493
                                                            1.00 41.08
                                  51.923
              OH2 WAT G 333
HETATM 6358
                                                   79.363
                                                            1.00 39.05
                                          27.611
              OH2 WAT G 334
                                  49.511
нетатм 6359
                                                            1.00 41.42
              OH2 WAT G 335
                                  69.385
                                           0.852 110.192
HETATM 6360
                                           2.479
                                                  101.880
                                                           1.00 42.50
                                  40.952
              OH2 WAT G 336
HETATM 6361
                                                            1.00 54.22
                                  32.998
                                           7.200 103.784
              OH2 WAT G 337
HETATM 6362
                                                            1.00 52.69
                                          15.261 136.205
                                  54.366
              OH2 WAT G 338
HETATM 6363
                                  35.674
                                          13.727
                                                   89.792
                                                            1.00 35.83
              OH2 WAT G 339
HETATM 6364
                                  66.606 -21.361
                                                   87.138
                                                            1.00 46.26
HETATM 6365
              OH2 WAT G 340
                                           4.708 131.550
                                                            1.00 45.27
              OH2 WAT G 341
                                  72.053
HETATM 6366
                                  28.072
                                          -1.358
                                                   70.419
                                                            1.00 34.92
              OH2 WAT G 342
HETATM 6367
                                                            1.00 52.99
                                          -3.981
                                                   76.422
                                  23.611
              OH2 WAT G 343
HETATM 6368
                                  53.684
                                           2.564 122.150
                                                            1.00 58.16
HETATM 6369
              OH2 WAT G 344
                                          -6:528 125.556
                                                            1.00 34.71
              OH2 WAT G 345
                                  30.624
HETATM 6370
                                          13.838 113.997
                                                            1.00 44.91
                                  27.870
              OH2 WAT G 346
HETATM 6371
                                          -9.588 116.327
                                                            1.00 55.34
                                  31.903
              OH2 WAT G 347
HETATM 6372
                                  71.763
                                                            1.00 48.99
                                          15.094
                                                   63.739
HETATM 6373
              OH2 WAT G
                        348
                                          -2.536 114.760
                                                            1.00 37.19
                                  25.258
              OH2 WAT G 349
HETATM 6374
                                                            1.00 42.32
                                  43.765
                                          12.162
                                                   78.143
              OH2 WAT G 350
HETATM 6375
                                                            1.00 33.70
                                  32.452
                                           5.338
                                                   73.909
              OH2 WAT G 351
HETATM 6376
                                          -5.770 101.894
                                                            1.00 46.40
                                  52.896
HETATM 6377
              OH2 WAT G 352
                                                            1.00 34.62
                                           4.242 115.852
                                  47.968
              OH2 WAT G 353
HETATM 6378
                                                            1.00 49.80
                                          -9.302
                                                   90.596
HETATM 6379
                        354
                                  38.561
              OH2 WAT G
                                                   74.354
                                                            1.00 56.40
                                  63.791
                                          17.454
HETATM 6380
              OH2 WAT G 355
                                           2.648 133.760
                                                            1.00 50.00
              OH2 WAT G 356
                                  41.360
HETATM 6381
                                                            1.00 38.01
                                           -7.937 122.328
              OH2 WAT G 357
                                  42.467
HETATM 6382
                                  50.890
                                          -0.362 116.668
                                                            1.00 39.26
              OH2 WAT G 358
HETATM 6383
                                         -23.881
                                                   67.865
                                                            1.00 55.18
              OH2 WAT G 359
                                  54.217
нетатм 6384
                                                            1.00 38.83
                                           9.539 105.032
              OH2 WAT G 360
                                  64.959
HETATM 6385
                                  58.113 -19.846
                                                   82.288
                                                            1.00 38.60
HETATM 6386
              OH2 WAT G 361
                                          -1.140
                                                   93.572
                                                            1.00 31.47
нетатм 6387
              OH2 WAT G
                        362
                                  42.245
                                                           1.00 54.89
                                          17.770 125.885
                                  73.552
              OH2 WAT G 363
HETATM 6388
                                  68.769
                                                            1.00 45.53
                                          15.898 106.810
HETATM 6389
              OH2 WAT G 364
                                  37.543
                                          19.031
                                                   78.866
                                                            1.00 45.15
HETATM 6390
              OH2 WAT G
                        365
                                                   95.087
                                                            1.00 44.99
                                            6.906
              OH2 WAT G 366
                                  55.583
HETATM 6391
                                                            1.00 36.58
                                  41.284
                                            9.699
                                                   78.250
HETATM 6392
              OH2 WAT G 367
                                            5.332 126.362
                                                            1.00 46.60
                                  25.203
HETATM 6393
              OH2 WAT G
                        368
                                                   95.104
                                                            1.00 47.85
                                          -5.006
HETATM 6394
              OH2 WAT G 369
                                  74.742
                                                            1.00 51.46
                                  70.349
                                          19.871
                                                    69.925
              OH2 WAT G 370
HETATM 6395
                                                    94.720
                                                            1.00 38.66
                                  42.93.6
                                          20.631
HETATM 6396
              OH2 WAT G
                        371
                                                            1.00 44.01
                                  34.162 -16.114 114.141
              OH2 WAT G 372
HETATM 6397
                                                            1.00 44.66
                                                  100.275
                                  33.863
                                          16.838
              OH2 WAT G 373
HETATM 6398
                                                   86.140
                                                            1.00 43.89
                                          12.569
                                  21.613
HETATM 6399
              OH2 WAT G
                        374
                                  35.751 -13.302 100.583
70.095 13.395 117.505
                                                            1.00 53:53
              OH2 WAT G 375
HETATM 6400
                                                            1.00 52.02
              OH2 WAT G 376
HETATM 6401
                                         19.108 131.799
                                                            1.00 46.47
                                  41.853
HETATM 6402
              OH2 WAT G
                        377
```

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HETATM 640	3 OH2 WAT G 378	55.780 -14.986 65.487 1.00 49.09
HETATM 640	4 OH2 WAT G 379	40.990 21.205 91.611 1.00 41.02
HETATM 640	5 OH2 WAT G 380	40 100 4 000 410 000 4 00 44 0.
HETATM 640	6 OH2 WAT G 381	48.157
HETATM 640	7 OH2 WAT G 382	30.221 27.743 109.194 1.00 39.92
HETATM 640	8 OH2 WAT G 383	30.221 27.743 109.194 1.00 39.92 49.926 -12.826 118.421 1.00 58.95
HETATM 640	9 OH2 WAT G 384	48.157
HETATM 641		58.226 -25.990 71.378 1.00 48.18
HETATM 641		40 405 17 044 129 741 1 00 42 02
HETATM 641		40.495 17.944 128.741 1.00 43.82
HETATM 6413		31.943 6.301 109.475 1.00 35.53 47.277 2.559 100.509 1.00 43.00
	_	47.277 2.559 100.509 1.00 43.00 38.862 9.112 102.620 1.00 31.70
HETATM 6414		
HETATM 6415		71.652 14.568 105.167 1.00 49.63
HETATM 6416	•	68.554 -10.518 73.331 1.00 38.16 70.496 -16.160 84.425 1.00 32.16
HETATM 6417	OH2 WAT G 392	70.496 -16.160 84.425 1.00 32.16
HETATM 6418	OH2 WAT G 393	44.698 -24.950 75.603 1.00 43.38 56.172 15.369 55.027 1.00 47.44
HETATM 6419	OH2 WAT G 394	56.172 15.369 55.027 1.00 47.44
HETATM 6420		46.150 -9.441 99.999 1.00 47.98
HETATM 6421	OH2 WAT G 396	46.150 -9.441 99.999 1.00 47.98 26.892 -8.356 89.057 1.00 34.99 31.737 14.380 90.395 1.00 50.78 36.261 -13.824 62.777 1.00 50.86 37.312 15.242 134.977 1.00 43.57 33.728 13.773 126.419 1.00 57.13
HETATM 6422	OH2 WAT G 397	31.737 14.380 90.395 1.00 50.78
HETATM 6423	OH2 WAT G 398	36.261 -13.824 62.777 1.00 50.86
HETATM 6424	OH2 WAT G 399	37.312 15.242 134.977 1.00 43.57
HETATM 6425	OH2 WAT G 400	33.728 13.773 126.419 1.00 57.13
HETATM 6426	OH2 WAT G 401	45.269 27.937 130.311 1.00 49.55
HETATM 6427	OH2 WAT G 402	44.887 -17.414 111.508 1.00 54.29
HETATM 6428	OH2 WAT G 403	36.261 -13.824 62.777 1.00 50.86 37.312 15.242 134.977 1.00 43.57 33.728 13.773 126.419 1.00 57.13 45.269 27.937 130.311 1.00 49.55 44.887 -17.414 111.508 1.00 54.29 68.928 0.455 136.711 1.00 49.90 43.271 -21.571 64.425 1.00 48.61 24.243 -4.781 108.590 1.00 51.05
HETATM 6429	OH2 WAT G 404	43.271 -21.571 64.425 1.00 48.61
HETATM 6430	OH2 WAT G 405	24.243 -4.781 108.590 1.00 51.05
HETATM 6431	OH2 WAT G 406	54.828 5.311 59.009 1.00 43.43
HETATM 6432	OH2 WAT G 407	53.460 27.992 124.076 1.00 47.83
HETATM 6433	OH2 WAT G 408	24.243 -4.781 108.590 1.00 51.05 54.828 5.311 59.009 1.00 43.43 53.460 27.992 124.076 1.00 47.83 70.833 -18.390 85.386 1.00 49.26 71.497 15.287 113.071 1.00 34.52 36.407 -18.480 110.466 1.00 55.43 26.220 -9.551 78.158 1.00 47.69 52.319 26.326 82.038 1.00 42.00
HETATM 6434	OH2 WAT G 409	71.497 15.287 113.071 1.00 34.52
HETATM 6435	OH2 WAT G 410	36.407 -18.480 110.466 1.00 55.43
HETATM 6436	OH2 WAT G 411	26.220 -9.551 78.158 1.00 47.69
HETATM 6437	OH2 WAT G 412	52.319 26.326 82.038 1.00 42.00
HETATM 6438	OH2 WAT G 413 OH2 WAT G 414	76.173 14.097 122.253 1.00 44.90 58.379 6.335 123.024 1.00 54.61
HETATM 6439	OH2 WAT G 414	58.379 6.335 123.024 1.00 54.61
HETATM 6440	OH2 WAT G 415	72.162 -16.705 82.719 1.00 50.63
HETATM 6441	OH2 WAT G 416	63.557 26.152 65.944 1.00 39.83
HETATM 6442	OH2 WAT G 417 OH2 WAT G 418	38.935 23.070 122.742 1.00 52.57
HETATM 6443	OH2 WAT G 418	55.256 -10.714 124.501 1.00 42.38 55.443 -9.037 110.170 1.00 46.47
HETATM 6444	OH2 WAT G 419	55.443 -9.037 110.170 1.00 46.47
HETATM 6445	OH2 WAT G 420	73.873 16.578 123.288 1.00 46.54
HETATM 6446	OH2 WAT G 421	74.426 12.663 117.527 1.00 43.62
HETATM 6447	OH2 WAT G 422	52.374 -0.368 51.502 1.00 56.99
HETATM 6448	OH2 WAT G 423	60.339 20.215 84.713 1.00 36.27
HETATM 6449	OH2 WAT G 424 OH2 WAT G 425	48.308 1.354 54.561 1.00 38.53
HETATM 6450	OH2 WAT G 425	61.757 21.606 115.976 1.00 61.09
HETATM 6451	OH2 WAT G 426	33.222 -14.916 119.528 1.00 51.12 47.477 3.359 112.298 1.00 46.10
HETATM 6452	OH2 WAT G 427	47.477 3.359 112.298 1.00 46.10
HETATM 6453	OH2 WAT G 428	39.909 2.272 138.388 1.00 35.33
HETATM 6454	OH2 WAT G 429	57.829 15.336 126.262 1.00 62.59
HETATM 6455	OH2 WAT G 430	48.917 -5.857 119.191 1.00 51.45
HETATM 6456	OH2 WAT G 431	44.139 -3.812 132.964 1.00 44.91
HETATM 6457	OH2 WAT G 432	38.885 18.594 95.398 1.00 50.23
HETATM 6458	OH2 WAT G 433	52.628 -7.064 55.271 1.00 38.96
HETATM 6459	OH2 WAT G 434	60.644 -0.731 101.129 1.00 47.30
HETATM 6460	OH2 WAT G 435	64.772 5.808 71.942 1.00 50.81
HETATM 6461	CH2 WAT G 436	39.571 16.705 80.180 1.00 34.07
HETATM 6462	OH2 WAT G 437	32.791 -0.551 65.371 1.00 41.40
HETATM 6463	OH2 WAT G 438	58.318 -7.989 60.087 1.00 46.94
HETATM 6464	OH2 WAT G 439	26.982 5.474 120.408 1.00 46.28
HETATM 6465	OH2 WAT G 440	72.138 1.233 90.050 1.00 50.13
HETATM 6466	OH2 WAT G 441	29.494 10.971 118.393 1.00 56.30
HETATM 6467	OH2 WAT G 442	69.232 5.594 113.941 1.00 58.17
HETATM 6468	OH2 WAT G 443	61.459 11.576 71.140 1.00 61.67
		• • • • • • • • • • • • • • • • • • • •

HETATM	6469	он2	WAT	G	444	59.592	2.195	58.518		42.66
HETATM	6470	OH2	_WAT	G	445	47.407	6.152	111.310		45.14
HETATM		OH2	WAT	G	446	36.254	18.203	99.930		44.76
HETATM		OH2	WAT	G	447	49.525	32.050	116.235	1.00	47.72
HETATM			TAW			21.801	-5.358	81.109	1.00	42.07
HETATM		OH2	WAT	G	449	52.131	-14.007	95.380	1.00	40.76
HETATM			WAT	-		39.712	-19.983	72.499	1.00	51.69
HETATM			WAT			67.651	5.620	67.102	1.00	42.38
HETATM			WAT			77.344	1.313	79.207	1.00	63.64
HETATM			WAT			55.249	-29.426	86.187	1.00	44.98
HETATM			WAT			64.429	-11.004	98.104	1.00	49.12
HETATM			WAT	-		45.456	-0.814	129.510	1.00	61.60
HETATM	-		WAT				-14.790	68.028	1.00	40.08
HETATM			TAW				5.611	94.924	1.00	58.32

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Figure 19-1

			٠		Residue	∄ X	Y	Z	B Segment	ID	
ATOM '	1				2	45.868	37.229	75.022	1.00 57.10	A.	AAA
ATOM	2		ALA		2	. 46.761	38.761	73.244	1.00 55.49		444
MOTA	3		ALA		2	46.339	39.800		1.00 55.57		AAA
ATOM	4 5		ALA ALA		2 2	48.280 47.062	37.746 37.537	74.937 74.110	1.00 57.26		AAA
MOTA MOTA	6		LYS		3	46.976	38.628	71.938	1.00 56.37 1.00 53.94		AAA AAA
ATOM	7				3	46.721	39.716	71.002	1.00 51.97		AAA
ATOM	. 8	CE			3	47.815	39.778	69.939	1.00 53.86		LAA
ATOM	9	CG			3	49.223	39.876	70.490	1.00 56.47		AA
ATOM	10	22	LYS	A	3	50.252	39.670	69.387	1.00 57.84		AA
ATOM	11		LYS		3	51.654	39.597	69.957	1.00 58.89	AA	AA
ATOM	12	ΝZ	LYS		3 -	52.643	39.283	68.895	1.00 59.33		AA
ATOM	13	C	LYS		3	45.393	39.494	70.305	1.00 49.57		AA
ATOM	14 15	0	LYS VAL		3 4	44.894 44.826	38.373 40.574	70.246 69.777	1.00 49.33 1.00 46.23		AA
ATOM ATOM	16	CA	AYT		4	43.561	40.516	69.056	1.00 40.23		AA AA
ATOM	17	СЗ	VAL		4	42.543	41.516	69.630	1.00 42.26		AA
ATOM	18		L VAL		4	41.213	41.352	68.940	1.00 41.51		AA
ATOM	19	CG	Z VAL	A	4	42.401	41.307	71.128	1.00 42.00		AA
ATOM	20	c	VAL		4	43.918	40.913	67.638	1.00 39.94		AA
MOTA	21	Ü	VAL		4	44.332	42.032	67.395	1.00 40.39	AA	Añ.
MOTA	22	N	LYS		5	43.766	40.001	66.695	1.00 36.94		ΑÀ
MOTA	23	CA	LYS LYS		5 5	44.142	40.305	65.323	1.00 34.10		ĀĀ
ATOM ATOM	24 25	CB	LYS		5	45.179 46.424	39.290 39.182	64.846 65.698	1.00 35.02 1.00 34.07		ÄÄ
ATOM	26	22	LYS		5	47.233	40.452	65.652	1.00 33.44		AA AA
ATOM	27	CΞ	LYS		5	48.565	40.239		1.00 32.38		ÀÀ
ATOM	28	::Z	LYS	A	5	49.372	41.460	66.222	1.00 31.26		ÀÀ
ATOM	29	С	LYS		5	42.997	40.293	64.333	1.00 31.38		AA
MOTA	30	0	LYS		5	42.053	39.523	64.466	1.00 31.74		AA
ATOM	31	11	LEU .		6	43.090	41.142	63.326	1.00 28.60		AA
ATOM	32	CA	LEU .		6 6	42.075	41.167 42.580	62.289	1.00 26.90	AA	
ATOM ATOM	33 34	CB	LEU .		6	41.530 40.321	42.748	62.067 61.129	1.00 26.43 1.00 25.89	AA AA	
MOTA	35	CD1			6	40.108	44.224	60.826	1.00 25.50		aa Aa
ATOM	36		LEU		6	40.550	42.032	59.828	1.00 26.64	λA	
ATOM	37	C	LEU Z		6	42.818	40.701	61.049	1.00 25.74	AA	
MOTA	38	2	LEU		6	43.877	41.226	60.717	1.00 24.60	AA	AA
ATOM	39	:1	ILE A	À	7	42.282	39.704	60.367	1.00 25.90	AA	
ATOM	40	CA.	ELE ?	y	7 7	42.939	39.212	59.173	1.00 26.75	λA	
MOTA	41 42	03 032	TLE A		7	42.839 43.474	37.712 37.227	59.089 57.783	1.00 26.58 1.00 27.88	AA	
atom atom	43	232	=== 2	Ä	÷	43.528	37.116	60.310	1.00 27.02	AA AA	
ATOM	44	CDI		A	7	43.507	35.640	60.350	1.00 27.46	AA	
ATOM	45	2	ILE ?	A	7	42.339	39.814	57.929	1.00 26.70	AA	
MOTA	46	J	ILE A		7	41.162	39.655	57.681	1.00 27.68	λA	
ATOM	47	N	GLY A		8	43.144	40.509	57.142	1.00 27.94	λA	
atom	-:8	CA	GLY A		8	42.598	41.110	55.944	1.00 29.78	AA	
ATOM	÷0	0	GLY A		8	43.587	41.789	55.027 55.264	1.00 30.38 1.00 29.39	AA	
ATOM	50 51		THR A		8 9	44.785 43.051	41.765 42.395	53.204	1.00 29.39	AA	
aton aton	51 52	N CA	THR A		9	43.832	43.106	52.962	1.00 32.41	AAI AAI	
ATCM	53	J3	THR A		9	44.606	42.112	52.064	1.00 31.12	AA	
ATOM	54		THR A		9	45.324	42.825	51.053	1.00 30.74	AA	
ATOM	55	CG2	THR A	A.	9	43.654	41.140	51.411	1.00 30.27	AA	AA
ATOM	56	C	THR A		9	42.886	43.939	52.091	1.00 32.94	AA	
ATOM	57	3	THR A		9	41.705	43.625	51.993	1.00 33.62	AAA	
ATCM	58	N	LEU A		10	43.396	45.009	51.485	1.00 33.20	AAA	
ATOM	59	CA	LEU A		10	42.573	45.840	50.611	1.00 33.29	AAA	
ATOM	50	CB	LEU A		10 10	43.117	47.275	50.484 51.666	1.30 33.12 1.00 32.95	AAA	
ATCM	51 62	CS CD1	LEU A		_0 10	43.142	48.245 48.386	52.288	1.00 32.95	ኣልሩ ኣልሩ	
atom atom	63		LEU A		10	44.126	47.734	52.675	1.00 34.71	AAA	
ATCM	64	c	LEU A		10	42.527	45.231	49.218	1.00 33.18	AAA	
ATCM	65	ō	LEU A		10	41.876	45.768	48.328	1.00 32.52	AAA	
ATCM		N	ASP A		11	43.230	44.121	49.022	1.00 33.56	AAA	

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> mo>f	67	CA	ASP	Δ	11	43.240	43.489	47.716	1.00 34.24	AAAA
MOTA			ASP		11	44.393	42.499	47.607	1.00 35.81	AAAA
MOTA	68	CB				45.739	43.190	47.604	1.00 37.57	AAAA
MOTA	69	CG	ASP		11				1.00 37.95	AAAA
ATOM	70		ASP		11	45.890	44.178	46.855		AAAA
MOTA	71	OD2	ASP	Α	11	46.650	42.750	48.332	1.00 40.31	
ATOM -	72	С	ASP	Α	11	41.929	42.813	47.341	1.00 34.03	AAAA
MOTA	73	0	ASP	Α	11	41.629	42.652	46.150	1.00 34.80	AAAA
MOTA	74	N	TYR		12	41.142	42.417	48.335	1.00 32.34	AAAA
	75	CA	TYR		12	39.871	41.803	48.017	1.00 32.53	AAAA
MOTA					12	39.043	41.569	49.290	1.00 31.32	AAAA
MOTA	76	CB	TYR			39.551	40.438	50.162	1.00 29.95	AAAA
MOTA	77	CG	TYR		12			51.469	1.00 28.52	AAAA
MOTA	78		TYR		12	39.983	40.669			AAAA
MOTA	79	CE1	TYR	Α	12	40.413	39.614	52.279-	1.00 28.03	
MOTA	80	CD2	TYR	Α	12	39.568	39.128	49.688	1.00 28.47	AAAA
ATOM	81	CE2	TYR	Α	12	39.992	38.083	50.483	1.00 28.47	AAAA
ATOM	82	CZ	TYR	Α	12	40.408	38.330	51.775	1.00 28.43	- AAAA
	83	OH	TYR		12	40.786	37.277	52.569	1.00 29.86	AAAA
MOTA	84	C	TYR		12	39.146	42.749	47.066	1.00 33.16	AAAA
ATOM					12	38.554	42.324	46.082	1.00 33.36	AAAA
ATOM	85	0	TYR			39.237	44.041	47.356	1.00 34.76	AAAA
MOTA	86	N	GLY		13		45.065	46.546	1.00 36.60	AAAA
ATOM	87	CA	GLY		13	38.594			1.00 37.85	AAAA
ATOM	88	С	GLY	A	13	38.814	44.961	45.052		AAAA
MOTA	89	0	GLY	Α	13	38.105	45.591	44.275	1.00 37.40	
MOTA	90	N	LYS	Α	14	39.799	44.171	44.647	1.00 39.55	AAAA
ATOM	91	CA	LYS	Α	14	40.091	43.981	43.231	1.00 40.66	AAAA
ATOM	92	CB	LYS		14	41.605	43.977	42.995	1.00 42.26	AAAA
	93	CG	LYS		14	42.300	45.309	43.239	1.00 44.54	AAAA
MOTA	94	CD	LYS		14	41.820	46.445	42.304	1.00 46.32	AAAA
ATOM			LYS		14	42.033	46.158	40.810	1.00 46.64	AAAA
ATOM	95	CE				41.133	45.086	40.256	1.00 47.23	AAAA
MOTA	96	NZ	LYS		14	39.499	42.675	42.707	1.00 40.35	AAAA
ATOM	97	С	LYS		14			41.511	1.00 39.97	AAAA
ATOM	98	0	LYS	A	14	39.593	42.377			AAAA
ATOM	99	N	TYR	Α	15	38.897	41.901	43.605	1.00 39.95	AAAA
ATOM	100	CA	TYR	Α	15 -	38.300	40.617	43.245	1.00 40.30	
ATOM	101	CB	TYR	Α	15	38.962	39.490	44.050	1.00 38.46	AAAA
ATOM	102	CG	TYR	Α	15	40.472	39.519	44.021	1.00 37.01	AAAA
	103		TYR		15	41.213	39.136	45.137	1.00 36.24	AAAA
ATOM	104		TYR		15	42.604	39.220	45.144	1.00 35.73	AAAA
ATOM		CD2	TYR		15	41.163	39.976	42.902	1.00 36.84	AAAA
MOTA	105				15	42.556	40.064	42.898	1.00 36.53	AAAA
ATOM	106		TYR			43.271	39.689	44.028	1.00 36.24	AAAA
MOTA	107	CZ	TYR		15	44.648	39.816	44.042	1.00 36.49	AAAA
ATOM	108	OH	TYR		15			43.556	1.00 41.98	AAAA
MOTA	109	С	TYR		15	36.802	40.647		1.00 42.59	AAAA
ATOM	110	0	TYR	Α	15	36.288	39.786	44.280	1.00 42.81	· AAAA
ATOM	111	N	ARG	A	16	36.101	41.638	43.014		
ATOM	112	CA	ARG	Α	16	34.670	41.753	43.257	1.00 43.47	AAAA
ATOM	113	CB	ARG	Α	16	34.205	43.197	43.111	1.00 45.27	AAAA
	114	CG	ARG		16	35.021	44.234	43.833	1.00 48.06	AAAA
MOTA	115	CD	ARG		16	34.891	44.196	45.339	1.00 49.63	AAAA
ATOM			ARG		16	35.632	45.322	45.905	1.00 51.65	AAAA
ATOM	116	NE			16	35 382	46.602	45.622	1.00 52.71	AAAA
MOTA	117	CZ	ARG	_		34 406	46 931	44 781	1.00 53.28	AAAA
MOTA	118		ARG		16		47.560	46.162	1.00 53.43	AAAA
MOTA	119		ARG		16	36.124		42.230	1.00 42.86	AAAA
ATOM	120	С	ARG	Α	16	33.913	40.929		1.00 41.83	AAAA
ATOM	121	0	ARG	Α	16	34.455	40.541	41.193	1.00 41.63	
ATOM	122	N	TYR	Α	17	32.651	40.668	42.523	1.00 42.42	AAAA
ATOM	123	CA	TYR		17	31.818	39.942	41.590	1.00 42.76	AAAA
	124	CB	TYR		17	30.675	39.254	42.333	1.00 40.11	AAAA
MOTA	125	CG	TYR		17	31.097	38.061	43.180	1.00 38.35	AAAA
MOTA			TYR		17	32.169	38.148	44.071	1.00 36.15	AAAA
MOTA	126	CDI	TIK	~		32.519	37.069	44.874	1.00 34.76	AAAA
ATOM	127		TYR		17		36.855	43.116	1.00 36.40	AAAA
ATOM	128	CD2			17	30.386		43.912	1.00 35.31	AAAA
MOTA	129	CE2	TYR		17	30.726	35.776		1.00 35.00	AAAA
MOTA	130	CZ	TYR	Α	17	31.792	35.887	44.790	1.00 33.30	AAAA
ATOM	131	OH	TYR	A	17	32.115	34.814	45.584	1.00 33.29	AAAA
ATOM	132	C	TYR		17	31.296	41.000	40.613	1.00 44.43	AAAA
A.O.		•								•

ATOM	133	0	TYR	A 17	31.346	42.194	40.905	1.00 44.68	AAAA
MOTA	134	N	PRO	A 18	30.799	40.574	39.440	1.00 45.95	AAAA
MOTA	135	CD	PRO	A 18	30.707	39.175	38.994	1.00 46.08	AAAA
MOTA	136	CA	PRO				38.402	1.00 47.24	AAAA
ATOM	137	CB	PRO			40.482	37.312	1.00 47.69	AAAA
MOTA	138	CG	PRO		30.876		37.511	1.00 46.79	AAAA
ATOM	139	C	PRO		29.129	42.390	38.834	1.00 48.98	AAAA
MOTA	140	0	PRO		28.298	42.020	39.660	1.00 49.11	. AAAA
MOTA	141	N	LYS		29.114	43.593	38.253	1.00 50.59	AAAA
MOTA	142	CA	LYS		28.125	44.654	38.519	1.00 52.10	AAAA
ATOM	143	CB	LYS		27.876	45.466	37.246	1.00 54.41	AAAA
ATOM	144	CG	LYS LYS		29.120 28.747	45.911 46.508	36.498 35.142	1.00 57.78 1.00 59.34	AAAA
MOTA	145 146	CD	LYS		29.978	46.774	34.288	1.00 59.34	AAAA AAAA
ATOM. ATOM	147	NZ	LYS		29.616	47.277	32.932	1.00 61.03	AAAA
ATOM	148	C	LYS		26.764	44.162	39.012	1.00 51.53	AAAA
ATOM	149	ō	LYS		26.281	44.556	40.071	1.00 51.54	AAAA
ATOM	150	Ň	ASN		26.146	43.314	38.203	1.00 50.13	AAAA
ATOM	151	CA	ASN		24.831	42.750	38.482	1.00 48.44	AAAA
ATOM	152	СВ	ASN		24.336	42.061	37.209	1.00 49.67	AAAA
MOTA	153	CG	ASN	A 20	25.389	41.132	36.613	1.00 51.61	AAAA
MOTA	154	OD1	ASN	A 20	25.677	40.064	37.154	1.00 51.70	AAAA
ATOM	155		ASN		25.998	41.562	35.509	1.00 53.00	AAAA
ATOM	156	C	ASN		24.789	41.765	39.649	1.00 45.57	AAAA
ATOM	157	0	ASN .		23.764	41.127	39.877	1.00 44.67	AAAA
ATOM	158	N	HIS.		25.883	41.662	40.398	1.00 42.71	AAAA
ATOM	159	CA CB	HIS .		25.958 27.216	40.709 39.857	41.506 41.353	1.00 40.69 1.00 40.16	AAAA
MOTA MOTA	160 161	CG	HIS .		27.186	38.587	42.140	1.00 39.93	AAAA AAAA
ATOM	162		HIS.		27.329	38.353	43.467	1.00 39.27	AAAA
ATOM	163		HIS		26.951	37.359	41.557	1.00 39.47	AAAA
ATOM	164		HIS .		26.948	36.425	42.493	1.00 39.36	AAAA
ATOM	165		HIS A		27.174	37.003	43.660	1.00 39.44	AAAA
MOTA	166	С	HIS A	A 21	25.974	41.349	42.892	1.00 38.93	AAAA
MOTA	167	0	HIS A		26.660	.42.338	43.116	1.00 38.78	AAAA
ATOM	168	N	PRO A		25.229	40.778	43.853	1.00 37.11	AAAA
MOTA	169	CD	PRO A		24.371	39.579	43.814	1.00 36.09	AAAA
ATOM	170	CA	PRO A		25.224	41.361	45.199	1.00 35.81	AAAA
ATOM	171 172	CB CG	PRO A		24.473 23.464	40.306 39.810	46.012 45.003	1.00 36.04 1.00 36.19	AAAA AAAA
ATOM	173	C	PRO A		26.638	41.637	45.751	1.00 34.39	AAAA
ATOM ATOM	174	õ	PRO A		26.867	42.653	46.417	1.00 34.09	AAAA
ATOM	175	N	LEU A		27.572	40.731	45.451	1.00 31.98	AAAA
ATOM	176	CA	LEU A		28.954	40.827	45.900	1.00 29.65	AAAA
ATOM	177	CB	LEU A	A 23	29.564	39.432	46.014	1.00 27.88	AAAA
MOTA	178	CG	LEU A	A 23	28.896	38.528	47.048	1.00 27.31	AAAA
MOTA	179		LEU A		29.656	37.217	47.149	1.00 26.64	TAAA
ATOM	180		LEU A		28.879	39.212	48.399	1.00 26.75	AAAA
ATOM	181	C	LEU A		29.838		45.018	1.00 29.20	AAAA
ATOM	182	0	LEU A		31.057	41.606 42.582	45.028	1.00 28.38	AAAA
MOTA	183		LYS A		29.204	42.562	44.259 43.389	1.00 29.27	AAAA AAAA
ATOM	184 185	CA CB	LYS A		29.903 28.881	44.091	42.405	1.00 29.75	AAAA
ATOM	186	CG	LYS A		29.328	45.265	41.601	1.00 32.55	AAAA
MOTA MOTA	187	CD	LYS A		28.537	46.526	41.994	1.00 34.37	AAAA
ATOM	188	CE	LYS A		27.025	46.337	41.835	1.00 34.32	AAAA
ATOM	189	NZ	LYS A		26.221	47.542	42.208	1.00 34.37	AAAA
ATOM	190	C	LYS A		30.580	44.620	44.224	1.00 28.14	AAAA
ATOM	191		LYS A		31.617	45.162	43.840	1.00 27.93	AAAA
ATOM	192	N	ILE A		29.990	44.919	45.377	1.06 27.07	· · AAAA
ATOM	193	CA	ILE A	. 25	30.468	45.945	46.296	1.00 25.82	AAAA
MOTA	194	CB	ILE A		29.425	46.262	47.364	1.00 25.37	AAAA
ATOM	195		ILE A		28.190	46.846	46.737	1.00 25.71	AAAA
MOTA	196		ILE A		29.142	44.979	48.157	1.00 25.26	AAAA
ATOM	197		ILE A		28.318	45.163	49.413	1.00 25.17	AAAA
MOTA	198	С	ILE A	25	31.700	45.550	47.095	1.00 25.28	, AAAA

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					32.037	44.379	47.183	1.00 24.48	AAAA
MOTA	199	0	ILE A			46.547	47.714	1.00 24.98	AAAA
MOTA	200	71	PRO A				47.638	1.00 24.98	AAAA
MOTA	201	CD	PRO A			47.980		1.00 24.44	AAAA
ATOM	202	CA	PRO A	26		46.367	48.543		AAAA
ATOM	203	CB	PRO A	A 26		47.792	48.701	1.00 24.75	
ATOM	204	CG	PRO A	26	33.435	48.546	47.538	1.00 25.51	AAAA
ATOM	205	C	PRO A		33.021	45.838	49.862	1.00 23.42	AAAA
	206	0	PRO A		31.930	46.233	50.272	1.00 22.12	AAAA
MOTA	207	N	ARG A		 :	44.960	50.532	1.00 23.06	AAAA
ATOM			ARG P			44.421	51.776	1.00 23.04	AAAA
MOTA	208	CA	ARG A			43.043	51.492	1.00 22.20	AAAA
MOTA	209	CB				43.152	50.503	1.00 19.84	AAAA
MOTA	210	CG	ARG A			41.844	50.160	1.00 18.64	AAAA
MOTA	211	CD	ARG A				51.315	1.00 16.51	AAAA
ATOM	212	NE	ARG A			41.168	51.982	1.00 16.57	AAAA
ATOM	213	cz	ARG A			40.188		1.00 17.50	AAAA
ATOM	214	NH1	ARG A	A 27		39.763	51.605		AAAA
ATOM	215	NH2	ARG A	A 27	30.185	39.643	53.038	1.00 16.45	
MOTA	216	C	ARG A	A 27	34.265	44.381	52.905	1.00 23.62	AAAA
MOTA	217	ō	ARG A		34.107	45.077	53.919	1.00 23.69	AAAA
	218	N	VAL A		35.305	43.570	52.736	1.00 24.25	AAAA
ATOM	219	CA	VAL			43.466	53.737	1.00 23.36	AAAA
ATOM			VAL A			42.062	53.671	1.00 22.75	AAAA
MOTA	220	CB				42.031	54.475	1.00 22.95	AAAA
MOTA	221		VAL A			41.011	54.249	1.00 22.20	AAAA
MOTA	222		VAL			44.609	53.511	1.00 23.70	AAAA
ATOM .	223	Ç	VAL			45.156	54.455	1.00 22.62	AAAA
MOTA	224	0	VAL			44.989	52.253	1.00 24.27	AAAA
MOTA	225	N	SER A	_			51.910	1.00 26.03	AAAA
MOTA	226	CA	SER A			46.082	50.381	1.00 25.95	AAAA
MOTA	227	CB	SER A			46.178		1.00 27.57	AAAA
ATOM	228	OG	SER .	A 29		46.417	49.716		AAAA
MOTA	229	С	SER A	A 29		47.354	52.440	1.00 25.52	AAAA
ATOM	230	0	SER 3	A 29		48.311	52.828	1.00 25.49	
MOTA	231	N	LEU .	A 30	36.468	47.342	52.448	1.00 26.09	AAAA
	232	CA	LEU .		35.692	48.471	52.926	1.00 26.39	AAAA
MOTA	233	CB	LEU			48.365	52.393	1.00 25.89	AAAA
ATOM	234	CG	LEU			49.470	52.755	1.00 27.15	AAAA
MOTA			LEU			49.101	53.999	1.00 26.34	AAAA
MOTA	235				·	50.813	52.897	1.00 25.81	AAAA
ATOM	236		LEU .			48.534	54.453	1.00 26.26	AAAA
MOTA	237	C	LEU .			49.612	55.037	1.00 27.50	AAAA
MOTA	238	0	LEU .			47.379	55.097	1.00 25.57	AAAA
MOTA	239	N	LEU	-		47.343	56.545	1.00 26.87	AAAA
MOTA	240	CA	LEU .				57.029	1.00 27.28	AAAA
MOTA	241	CB	LEU .			45.900	58.383	1.00 27.87	AAAA
MOTA	242	CG	LEU .			45.563			AAAA
ATOM	243	CD1	LEU	A 3:	35.855	44.313	58.906	1.00 27.01	AAAA
MOTA	244	CD2	LEU .	A 3	L 35 261	46.706	59.372	1.00 26.32	
MOTA	245	С	LEU .	A 3:	37.087	48.003	57.012	1.00 28.08	AAAA
	246	ō	LEU		37 094	48.854	57.901	1.00 27.42	AAAA
MOTA	247	Ŋ	LEU			47.584	56.409	1.00 29.52	AAAA
MOTA		CA	LEU			48.121	56.750	1.00 30.96	AAAA
MOTA	248		LEU			47.394	55.950	1.00 31.58	AAAA
ATOM	249	CB				45.904	56.293	1.00 31.63	AAAA
MOTA	250	CG	LEU			45.246	55.380	1.00 31.31	AAAA
ATOM	251		LEU		•	45.780	57.743	1.00 32.23	AAAA
ATOM	252	CD2	LEU			49.635	56.543	1.00 31.59	AAAA
ATOM	253	С	LEU				57.416	1.00 31.70	AAAA
ATOM	254	0	LEU	A 3:		50.345		1.00 32.72	AAAA
ATOM	255	N	ARG	A 3		50.129	55.398		AAAA
ATOM	256	CA	ARG	A 3		51.564	55.141	1.00 33.91	
	257	CB	ARG	_	38.643	51.903	53.743	1.00 35.10	AAAA
ATOM	258	CG	ARG	_		51.609	52.621	1.00 37.84	AAAA
MOTA		CD	ARG	_		52.412	51.374	1.00 39.33	AAAA
ATOM	259		ARG			51.806	50.580	1.00 42.51	AAAA
atom	260	NE	ARG			52.395	49.541	1.00 44.22	AAAA
MOTA	261	CZ	ALC:			53.617	49.163	1.00 44.61	AAAA
ATOM	262	NH1	ARG	2 3		51.752	48.861	1.00 45.23	AAAA
ATOM	263		ARG	A 3		52.305	56.168	1.00 33.48	AAAA
ATCM	264	С	ARG	A 3	3 30.334	26.303			•

		_			30	E2 200	F.C. C0.0	1.00 33.61	AAAA
ATOM	265	0	ARG A	33	38.713	53.390			
MOTA	266	N	PHE A	34	37.247	51.682	56.562	1.00 33.78	AAAA
ATOM	267	CA	PHE A	34	36.292	52.233	57.517	1.00 33.79	AAAA
	268	CB	PHE A	34	35.065	51.310	57.573	1.00 33.88	AAAA
MOTA	_						58.405	1.00 33.16	AAAA
MOTA	269	CG	PHE A	34	33.925	51.840			
MOTA	270	CD1	PHE A	34	33.108	52.856	57.925	1.00 32.77	AAAA
ATOM	271	CD2	PHE A	34	33.668	51.315	59.672	1.00 33.05	AAAA
	-		PHE A	34	32.044	53.343	58.695	1.00 32.86	AAAA
MOTA	272							1.00 33.07	AAAA
MOTA	273	CEZ	PHE A	34	32.607	51.797	60.454		
MOTA	274	CZ	PHE A	34	31.794	52.809	59.966	1.00 32.58	AAAA
	275	С	PHE A	34	. 36.881	52.414	58.918	1.00 34.01	. AAAA
MOTA					36.903	53.524	59.455	1.00 33.49	AAAA
MOTA	276	0	PHE A	34					•
ATOM	277	N	LYS A	35	37.350	51.324	59.516	1.00 34.00	AAAA
MOTA	278	CA	LYS A	35	37.928	51.401	60.843	1.00 33.90	AAAA
	279	CB	LYS A	35	38.230	50.010	61.362	1.00 34.07	AAAA
MOTA					37.000	49.190	61.662	1.00 33.94	AAAA
MOTA	280	CG	LYS A	35	_				
MOTA	281	CD	LYS A	35	37.414	47.810	62.106	1.00 35.31	AAAA
MOTA	282	CE	LYS A	35	38.062	47.072	60.948	1.00 35.91	AAAA
	283	NZ	LYS A	35	39.058	47.928	60.236	1.00 36.19	AAAA
MOTA					39.185	52.255	60.881	1.00 34.19	AAAA
MOTA	284	C	LYS A	35					
MOTA	285	0	LYS A	35	39.554	52.775	61.929	1.00 34.32	AAAA
ATOM	286	N	ASP A	36	39.853	52.384	59.745	1.00 33.99	AAAA
			ASP A	36	41.034	53.216	59.680	1.00 35.17	AAAA
ATOM	287	CA					58.388	1.00 37.40	AAAA
MOTA	288	CB	ASP A	36	41.812	52.943			
MOTA	289	CG	ASP A	36	42.964	53.908	58.186	1.00 38.64	AAAA
ATOM	290	OD1	ASP A	36	43.648	54.218	59.184	1.00 40.02	AAAA
	291		ASP A	36	43.201	54.341	57.035	1.00 38.74	AAAA
ATOM						54.670	59.724	1.00 35.85	AAAA
MOTA	292	С	ASP A	36	40.568				AAAA
ATOM	293	0	ASP A	36	41.231	55.527	60.306	1.00 36.88	
ATOM-	294	N	ALA A	37	39.420	54.940	59.111	1.00 34.96	AAAA
	295	CA	ALA A	37	38.851	56.280	59.108	1.00 34.47	AAAA
ATOM					37.751	56.373	58.067	1.00 33.80	AAAA
MOTA	296	CB	ALA A	37					AAAA
ATOM	297	С	ALA A	37	38.291	56.617	60.499	1.00 34.66	
ATOM	298	Q	ALA A	37	38.268	57.779	60.899	1.00 34.55	AAAA
	299	Ň	MET A	3.8	37.830	55.600	61.226	1.00 34.24	AAAA
MOTA					37.287	55.794	62.572	1.00 33.07	AAAA
MOTA	300	CA	MET A	38	-			1.00 32.82	AAAA
MOTA	301	CB	MET A	38	36.289	54.687	62.917		
ATOM	302	CG	MET A	38	35.084	54.559	61.996	1.00 32.72	AAAA
ATOM	303	SD	MET A	38	33.980	55.948	62.101	1.00 33.65	AAAA
			MET A	38	33.550	55.878	63.849	1.00 33.77	AAAA
ATOM	. 304	CE				55.724	63.583	1.00 33.12	AAAA
ATOM .	305	C	MET A	38	38.430				
MOTA	306	0	MET A	38	38.226	55.930	64.777	1.00 32.82	AAAA
MOTA	307	N	ASN A	39	39.628	55.428	63.090	1.00 32.64	AAAA
	308	CA	ASN A	39	40.805	55.266	63.935	1.00 32.38	AAAA
ATOM						56.600	64.589	1.00 32.93	AAAA
MOTA	309	CB	ASN A	39	41.200				AAAA
MOTA	310	CG	ASN A	39	41.393	57.736	63.571	1.00 34.40	
ATOM	311	OD1	ASN A	39	42.180	57.624	62.630	1.00 34.98	AAAA
	312		ASN A	39	40.677	58.838	63.772	1.00 33.52	AAAA
MOTA						54.212	65.009	1.00 31.69	AAAA
MOTA	313	С	ASN A	39	40.483			1.00 31.12	AAAA
MOTA	314	0	ASN A	39	·40.565	54.490	66.205		
ATOM	315	N	LEU A	40	40.095	53.010	64.570	1.00 31.76	AAAA
	316		LEU A	40	39.750	51.898	65.474	1.00 32.48	AAAA
MOTA		CA				51.559	65.386	1.00 32.55	AAAA
MOTA	317	CB	LEU A	40	38.259	31.333	65.300		
MOTA	.318	ÇG	LEU A	40	37.231	52.581	65.879	1.00 32.84	AAAA
ATOM	319		LEU A	40	35.837	52.089	65.554	1.00 33.79	AAAA
				40	37.372	52.798	67.376	1.00 32.45	AAAA
MOTA	320		LEU A			50.628	65.187	1.00 32.92	AAAA
MOTA	32,1	С	LEU A	40	40.555				
MOTA	322	0	LEU A	40	40.196	49.530	65.618	1.00 31.64	AAAA
	323	N	ILE A	41	41.652	50.794	64.464	1.00 34.12	AAAA
MOTA					42.508	49.680	64.116	1.00 36.07	AAAA
ATOM	324	CA	ILE A	41			62.811	1.00 35.51	AAAA
MOTA	325	CB	ILE A	41	42.017	48.991		1.00 33.37	AAAA
ATOM	326	CG2	ILE A	41	42.070	49.952	61.636	T.00 33.3/	
	327		ILE A	41	42.898	47.790	62.480	1.00 35.97	AAAA
MOTA				41	42.854	46.701	63.500	1.00 37.19	AAAA
MOTA	328		ILE A				63.916	1.00 38.85	AAAA
ATOM	329	С	ILE A	41	43.921	50.226		1.00 38.98	
ATOM	330	0	ILE A	41	44.106	51.346	63.413	1.00 20.38	AAAA
			_				-		•

ATOM	331	N	ASP A	42		44.914	49.446	64.329	1.00 40.61	AAAA
	332	CA	ASP A	42		46.309	49.843	64.181	1.00 42.57	AAAA
ATOM										
ATOM	333	CB	ASP A	42		46.973	50.021	65.553	1.00 42.42	AAAA
ATOM	334	CG	ASP A	42		46.316	51.110	66.381	1.00 42.27	AAAA
							52.250	65.883	1.00 41.20	AAAA
ATOM	335	ODI	ASP A	42		46.227				
ATOM_	336	OD2	ASP A	42		45.891	50.833	67.526	1.00 43.36	AAAA
						47.011	48.752	63.392	1.00 44.05	AAAA
ATOM	337	С	ASP A	42						
MOTA	338	0	ASP A	42		46.525	47.620	63.333	1.00 44.88	AAAA
						48.147	49.090	62.789	1.00 45.10	AAAA
MOTA	339	N	GLU A	-						
ATOM	340	CA	_GLU A	43		48.905	48.141	61.980	1.00 46.11	AAAA
	341	CB	GLU A	43		50.172	48.796	61.454	1.00 46.89	AAAA
MOTA									1.00 49.30	
ATOM	342	CG	GLU A	43		49.924	50.057	60.668		AAAA
MOTA	343	CD	GLU A	43	•	51.187	50.580	60.028	1.00 49.67	AAAA
						51.760	49.839	59.201	1.00 50.60	AAAA
ATOM	344		GLU A	43						
ATOM	345	OE2	GLU A	43		51.601	51.714	60.349	1.00 49.60	AAAA
	346	С	GLU A	43		49.290	46.859	62.701	1.00 46.27	- AAAA
MOTA										
ATOM	347	0	GLU A	43		49.214	45.773	62.131	1.00 46.00	AAAA
ATOM	348	N	LYS A	44		49.708	46.986	63.954	1.00 46.52	AAAA
								64.730	1.00 46.31	AAAA
MOTA	349	CA	LYS A	44		50.135	45.832			
ATOM	350	CB	LYS A	44		50.762	46.306	66.048	1.00 48.16	AAAA
		CG	LYS A	44		51.977	47.215	65.799	1.00 51.59	AAAA
MOTA	351									
MOTA	352	CD	LYS A	44		52.641	47.734	67.071	1.00 52.87	AAAA
ATOM	353	CE	LYS A	44		53.851	48.601	66.727	1.00 53.34	AAAA
								67.936	1.00 53.45	AAAA
MOTA	354	ΝZ	LYS A	44		54.615	49.033			
ATOM	355	С	LYS A	44		49.029	44.828	64.996	1.00 44.74	AAAA
		ō	LYS A	44		49.296	43.735	65.480	1.00 45.35	AAAA
ATOM	356									
ATOM	357	N	GLU A	45		47.793	45.190	64.659	1.00 42.49	AAAA
ATOM	358	CA	GLU A	45		46.638	44.320	64.894	1.00 40.54	AAAA
							45.125	65.517	1.00 40.55	AAAA
MOTA	359	CB	GLU A	45		45.493				
ATOM	360	CG	GLU A	45		45.788	45.731	66.882	1.00 38.87	AAAA
	361	CD	GLU A	45		44.663	46.618	67.360	1.00 37.57	AAAA
ATOM										AAAA
MOTA	362	OE1	GLU A	45		44.383	47.631	66.693	1.00 36.29	
ATOM	363	OE2	GLU A	45		44.056	46.300	68.399	1.00 38.44	AAAA
						46.126	43.648	63.630	1.00 39.15	AAAA
ATOM	364	С	GLU A	45						
MOTA	365	0	GLU A	45		45.301	42.737	63.681	1.00 39.29	AAAA
	366	N	LEU A	46		46.619	44.115	62.497	1.00 37.62	AAAA
MOTA							43.589	61.211	1.00 35.88	AAAA
ATOM	367	CA	LEU A	46		46.219				
MOTA	368	CB	LEU A	46		46.125	44.750	60.229	1.00 36.09	AAAA
	369	CG	LEU A	46		45.608	44.550	58.817	1.00 36.50	AAAA
MOTA									1.00 36.66	AAAA
ATOM	370	CD1	LEU A	46		44.182	44.021	58.843		
ATOM	371	CD2	LEU A	46		45.646	45.893	58.113	1.00 35.85	AAAA
			LEU A	46		47.211	42.542	60.714	1.00 34.97	AAAA
MOTA	372	С								
MOTA	373	0	LEU A	46		48.424	42.670	60.900	1.00 35.72	AAAA
ATOM	374	N	ILE A	47		46.680	41.484	60.118	1.00 33.25	AAAA
						47.497	40.411	59.560	1.00 30.92	AAAA
ATOM	375	CA	ILE A	47					-	
ATOM	376	CB	ILE A	47		47.144	39.024	60.167	1.00 31.22	AAAA
	377	CG2	ILE A	47		48.093	37.97C	59.640	1.00 28.55	AAAA
ATOM									1.00 32.04	AAAA
ATOM	378	CG1	ILE A	47		47.220	39.063	61.694		
ATOM	379	CD1	ILE A	47		48.596	39.241	62.242	1.00 34.13	AAAA
			ILE A	47		47.138	40.381	58.076	1.00 29.70	AAAA
MOTA	380	С								
ATOM	381	0	ILE A	47		45.956	40.373	57.714	1.00 28.42	AAAA
	382	N	LYS A	48		48.150	40.380	57.221	1.00 28.78	AAAA
ATOM						47.920	40.349	55.784	1.00 28.42	AAAA
ATOM	383	CA	LYS A	48						
ATOM	384	CB	LYS A	48		49.203	40.727	55.055	1.00 27.53	AAAA
				48		49.116	40.695	53.556	1.00 28.97	AAAA
MOTA	385	CG	LYS A							AAAA
MOTA	386	CD	LYS A	48		50.464	41.104	52.941	1.00 29.67	
ATOM	387	CE	LYS A	48		50.493	40.893	51.432	1.00 29.41	AAAA
				48		49.409	41.645	50.764	1.00 29.68	AAAA
ATOM	388	NZ	LYS A							
MOTA	389	C	LYS A	48		47.449	38.950	55.375	1.00 27.81	AAAA
		ō	LYS A	48		48.024	37.938	55.787	1.00 27.96	AAAA
ATOM	390								1.00 26.82	AAAA
ATOM	391	И	SER A	49		46.385	38.892	54.581		
	392	CA	SER A	49		45.854	37.611	54.141	1.00 26.41	AAAA
ATOM						44.514	37.795	53.420	1.00 25.40	AAAA
ATOM	393	CB	SER A	49						
ATOM	394	OG	SER A	. 49		43.541	38.349	54.276	1.00 25.58	AAAA
	395	C	SER A	49		46.814	36.891	53.207	1.00 26.03	AAAA
ATOM						47.462	37.513	52.373	1.00 26.98	AAAA
ATOM	396	0	SER A	49		21.204	٠,٠٠٠	-		
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MOTA	397	N	ARC	a a	50	46.910	35.576	53.354	1.00 25.51	AAAA
MOTA	398	CA	ARG	a a	50	47.755	34.794	52.474	1.00 25.45	AAAA
ATOM	399	CB	ARG		50	48.807	33.985	53.252	1.00 25.85	AAAA
ATOM	400	CG	ARG		50	48.229	32.819	54.009	1.00 27.16	AAAA
MOTA	401	CD	ARC		50	49.280	31.995	54.720	1.00 27.57	AAAA
MOTA	402	NE	ARG		50	48.673	30.896	55.482	1.00 27.90	AAAA
MOTA	403	CZ	ARG		50	48.106	29.820	54.946	1.00 28.34	AAAA
MOTA	404		ARG		50	48.055	29.672	53.630	1.00 28.19	AAAA
	405	NH2			50	47.592	28.884	55.735	1.00 28.62	AAAA
MOTA	406	C	ARG		50	46.806	33.834	51.762	1.00 24.91	AAAA
MOTA	407	õ	ARG		50	45.740	33.510	52.283	1.00 23.57	AAAA
ATOM		N	PRO		51	47.172	33.392	50.549	1.00 24.28	AAAA
MOTA	408	CD	PRO		51	48.361	33.761	49.770	1.00 24.13	AAAA .
MOTA	409		PRO		51	46.355		49.776	1.00 24.18	AAAA
MOTA	410	CA				47.012	32.512	48.390	1.00 24.24	AAAA
MOTA	411	CB	PRO		51	47.766	33.862	48.405	1.00 24.11	AAAA
ATOM	412	CG	PRO		51	46.473	31.070	50.393	1.00 23.69	AAAA
MOTA	413	C	PRO		51		30.680	50.839	1.00 24.13	AAAA
MOTA	414	0	PRO		51	47.545				
MOTA	415	N	ALA		52	45.381	30.325	50.422	1.00 23.36	AAAA
MOTA	416	CA	ALA		52	45.419	28.972	50.952	1.00 23.64	AAAA
MOTA	417	CB	ALA		52	44.012	28.405	51.029	1.00 23.86	AAAA
ATOM	418	C	ALA		52	. 46.260	28.145	49.994	1.00 23.58	AAAA
MOTA	419	0	ALA		52	46.240	28.383	48.806	1.00 24.52	AAAA
MOTA	420	N	THR		53	47.009	27.185	50.501	1.00 24.41	AAAA
MOTA	421	CA	THR		53	47.815	26.352	49.628	1.00 26.26	AAAA
MOTA	422	CB	THR		53	48.933	25.642	50.405	1.00 26.37	AAAA
MOTA	423		THR		53	48.355	24.763	51.375	1.00 26.51	AAAA
MOTA	424		THR		53	49.810	26.648	51.106	1.00 24.48	AAAA
MOTA	425	С	THR		53	46.889	25.299	49.034	1.00 27.63	AAAA
ATOM	426	0	THR		53	45.870	24.982	49.620	1.00 29.22	AAAA
ATOM	427	N	LYS		54	47.240		. 47.867	1.00 29.31	AAAA
MOTA	428	CA	LYS		54	46.450	23.752	47.189	1.00 30.61	AAAA
ATOM	429	CB	LYS		54	47.249	23.182		1.00 31.68	AAAA
ATOM	430	CG	LYS		54		22:020	45.304	1.00 34.38	AAAA
MOTA	431	CD	LYS		54	45.449	22.464	44.417	1.00 36.00	AAAA
ATOM	432	CE	LYS		54	45.943	22.850	43.025	1.00 37.55	AAAA
MOTA	433	NZ	LYS		54	46.425	21.664	42.236	1.00 37.57	AAAA
ATOM	434	C	LYS		54	46.127	22.640	48.170 48.176	1.00 31.26 1.00 31.72	AAAA AAAA
MOTA	435	0	LYS		54	45.025	22.097	49.006	1.00 31.72	AAAA
MOTA	436	N	GLU		55	47.102	22.312 21.260	50.011	1.00 32.29	AAAA
ATOM	437	CA	GLU		55	46.961 48.266	21.280	50.778	1.00 34.43	AAAA
MOTA	438	CB	GLU		55 55	48.265	19.901	51.706	1.00 38.39	AAAA
MOTA	439	CG	GLU GLU		55	49.513	19.839	52.584	1.00 41.46	AAAA
MOTA	440	CD	GLU		55	49.745	18.770	53.200	1.00 43.30	· AAAA
ATOM	441		GLU		55	50.245	20.859	52.672	1.00 42.45	AAAA
MOTA	442				55	45.851	21.555	51.013	1.00 30.43	AAAA
MOTA	443	C	GLU			45.048	20.681	51.332	1.00 30.59	AAAA
MOTA	444	0	GLU		55		22.782	51.517	1.00 28.23	AAAA
ATOM	445	N	GLU		56	45.822		52.488	1.00 27.69	AAAA
ATOM	446	CA	GLU		56	44.812	23.164	52.989	1.00 27.09	AAAA
MOTA	147	CB	GLU		56	45.078	24.588		1.00 26.64	
ATOM	448	CG	GLU		56	46.434	24.721	53.670 54.098	1.00 26.35	AAAA AAAA
MOTA	449	CD	GLU		56	46.769	26.135 27.057	53.265	1.00 25.12	AAAA
ATOM	450		GLU		56	46.615		55.255	1.00 25.70	AAAA
ATOM	451		GLU		56	47.213	26.315	51.914	1.00 26.99	AAAA
ATOM	452	C	GLU		56	43.408	23.043		1.00 26.25	
MOTA	453	0	GLU		56	42.495	22.574	52.588 50.659	1.00 26.25	AAAA AAAA
MOTA	454	N	LEU		57	43.252	23.447		1.00 27.26	
ATOM	455		LEU		57	41.965	23.389	49.967	1.00 27.17	AAAA
ATOM	456	CB	LEU		57	42.077	24.063	48.596		AAAA
ATOM	457		LEU		57	42.491	25.545	48.656	1.00 27.64	AAAA
MOTA	458		LEU		57	42.770	26.108	47.269	1.00 26.66 1.00 26.92	AAAA AAAA
MOTA	159		LEU		57	41.389	26.341	49.349	1.00 26.92	AAAA
ATOM	460	C	LEU		57	41.552	21.946	49.796	1.00 27.28	AAAA
MOTA	461	0	LEU		57	40.363	21.612	49.816 49.641	1.00 27.42	AAAA
MOTA	. 462	N	LEU	A	58	42.547	21.085	47.041	1.00 27.32	, and the second
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Figure 19-8

						44 455	40 455		
ATOM	463	CA	LEU A	58	42.293	19.675	49.457	1.00 26.10	AAAA
MOTA	464	CB	LEU A	58	43.486	19.019	48.794	1.00 25.43	AAAA
	465	CG	LEU A	58	43.623	19.577	47.385	1.00 26.66	AAAA
MOTA									
ATOM	466	CD1	. LEU A	58	44.760	18.884	46.705	1.00 27.12	AAAA
MOTA	467	CD2	LEU A	58	42.334	19.355	46.600	1.00 26.43	AAAA
	468	С	LEU A	58	41.938	18.956	50.731	1.00 25.79	AAAA
ATOM	_								
ATOM	469	0	LEU A	58	41.648	17.763	50.692	1.00 26.50	AAAA
ATOM	470	N	LEU A	59	41.977	19.666	51.858	1.00 24.91	AAAA
	471	CA	LEU A	59	41.595	19.070	53.136	1.00 25.15	AAAA
MOTA									
ATOM	472	CB	LEU A	59	41.958	19.991	54.322	1.00 25.44	AAAA
MOTA	473	CG	LEU A	59	43.423	20.280	54.710	1.00 24.67	AAAA
	474		LEU A	59	43.502	21.461	55.652	1.00 23.70	AAAA
ATOM									
MOTA	475	CD2	LEU A	59	44.044	19.044	55.357	1.00 24.08	AAAA
ATOM	476	С	LEU A	59	40.074	18.870	53.090	1.00 25.41	AAAA
ATOM	477	0	LEU A	59	39.503	18.266	53.993	1.00 25.88	AAAA
					39.436	19.392	52.031	1.00 25.05	AAAA
MOTA	478	N	PHE A	60					
ATOM	479	CA	PHE A	60	37.983	19.276	51.823	1.00 24.11	AAAA
ATOM	480	CB	PHE A	60	37.250	20.476	52.440	1.00 21.80	. AAAA
				60	35.778	20.534	52.098	1.00 20.07	AAAA
MOTA	481	CG	PHE A						
ATOM	482	CD1	PHE A	60	34.917	19.501	52.462	1.00 19.27	AAAA
ATOM	483	CD2	PHE A	60	35.249	21.628	51.399	1.00 19.82	AAAA
	484		PHE A	60	33.550	19.557	52.136	1.00 19.26	AAAA
ATOM									
MOTA	485	CE2	PHE A	60	33.890	21.688	51.071	1.00 17.45	AAAA
MOTA	486	CZ	PHE A	60	33.042	20.652	51.440	1.00 17.92	AAAA
	487	С	PHE A	60	37.557	19.139	50.345	1.00 24.02	AAAA
ATOM					36.846		49.974	1.00 23.27	AAAA
MOTA	488	0	PHE A	60		18.201			
ATOM	489	N	HIS A	61	37.982	20.079	49.511	1.00 24.40	AAAA
ATOM	490	CA	HIS A	61	37.626	20.053	48.099	1.00 25.04	AAAA
			HIS A		37.768	21.449	47.494	1.00 24.19	AAAA
ATOM	491	CB		61					
MOTA	492	CG	HIS A	61	36.744	22.429	47.979	1.00 24.44	AAAA
ATOM	493	CD2	HIS A	61	35.429	22.559	47.683	1.00 24.12	AAAA
	494		HIS A	61	37.038	23.444	48.864	1.00 24.36	AAAA
ATOM								1.00 23.18	AAAA
MOTA	495		HIS A	61	35.952	24.159.			
MOTA	496	NE2	HIS A	61	34.962	23.643	48.385	1.00 23.91	AAAA
ATOM	497	С	HIS A	61	38.416	19.054	47.253	1.00 25.60	AAAA
	498	ō	HIS A	61	39.596	18.805	47.498	1.00 26.94	AAAA
MOTA							46.244	1.00 26.68	AAAA
ATOM	499	N	THR A	62	37.754	18.496			
ATOM	500	CA	THR A	62	38.369	17.522	45.333	1.00 28.17	AAAA
ATOM	501	CB	THR A	62	37.290	16.695	44.614	1.00 28.15	AAAA
				62	36.544	17.541	43.731	1.00 28.10	AAAA
ATOM	502	OG1							
ATOM	503	CG2	THR A	62	36.334	16.094	45.629	1.00 28.24	AAAA
MOTA	504	С	THR A	62	39.226	18.217	44.278	1.00 29.28	AAAA
-	505	Ō	THR A	62	38.876	19.286	43.792	1.00 29.52	AAAA
ATOM						17.606	43.912	1.00 31.33	AAAA
ATOM	506	N	GLU A	63	40.344				
ATOM	507	CA	GLU A	63	41.249	18.202	42.928	1.00 32.42	AAAA
ATOM	508	CB	GLU A	63	42.333	17.219	42.536	1 00 34.37	AAAA
	509	CG	GLU A	63	43.304	16.869	43.609	1 00 37.20	AAAA
ATOM									
ATOM	510	CD	GLU A	63	44.427	16.022	43.052		AAAA
ATOM	511	OE1	GLU A	63	45.100	16.499	42.097	1.00 37.96	AAAA
	512	OE2		63	44.619	14.892	43.564	1.00 39.68	AAAA
ATOM							41.639	1.00 31.96	AAAA
MOTA	513	С	GLU A	63	40.607	18.687			
ATOM	514	0	GLU A	63	40.824	19.816	41.215	1.00 32.10	AAAA
MOTA	515	N	ASP A	64	39.845	17.814	40.998	1.00 31.52	AAAA
					39.204	18.165	39.753	1.00 31.36	AAAA
MOTA	516	CA	ASP A	64					
MOTA	517	CB	ASP A	64	38.301	17.018	39.295	1.00 33.99	AAAA
ATOM	518	CG	ASP A	64	37.213	16.694	40.302	1.00 37.38	AAAA
			ASP A	64	36.375	15.801	40.027	1.00 39.80	AAAA
ATOM	519					17.332	41.374	1.00 38.67	AAAA
MOTA	520	OD2	ASP A	64	37.188				
ATOM	521	С	ASP A	64	38.412	19.465	39.902	1.00 30.02	AAAA
	522	ō	ASP A	64	38.462	20.331	39.026	1.00 30.47	AAAA
ATOM							41.012	1.00 27.51	AAAA
ATOM	523	N	TYR A	65	37.695	19.608			
ATCM	524	CA	TYR A	65	36.918	20.814	41.248	1.00 26.03	AAAA
ATOM	525	CB	TYR A	65	36.010	20.654	42.467	1.00 25.42	AAAA
				65	35.339	21.946	42.866	1.00 24.90	AAAA
MOTA	526.	CG	TYR A			22.636	41.964	1,00 25.04	AAAA
- 501/				h h	34.525	22.b30	セエ・フロリ	1.00 43.04	AAAA
ATOM	527	CD1	TYR A	65					
MOTA ATOM	527 528	CD1 CE1	TYR A	65	33.914	23.823	42.308	1.00 25.01	AAAA

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MOTA	529	CD2	YYI	RА	65		35.525	22.486	44.136	1.00 24.65	AAAA
ATOM ·	530	CE		R· A			34.920	23.677		1.00 25.86	AAAA
										1.00 26.69	
MOTA	531	CZ		R A			34.110	24.349			AAAA
MOTA	532	OH		R A			33.499	25.543	43.924	1.00 27.20	Aaaa
MOTA	533	С	TY	RA	65		37.814	22.022	41.464	1.00 24.91	AAAA
MOTA	534	0	TY	R A	65	•	37.460	23.129	41.096	1.00 25.62	AAAA
ATOM	535	N	ILE	3 A	56		38.965	21.812	42.080	1.00 23.20	-AAAA
	536	CA	•	5 A	66	•	39.877	22.902	42.328	1.00 22,33	AAAA
MOTA								_			
MOTA	537			ΞA	66		40.924	22.520	43.402	1.00 21.45	AAAA
MOTA	538		ILE		66		41.927	23.652	43.617	1.00 20.00	AAAA
MOTA	539	CG1	. ILE	ΕΑ	66	-	40.220	22.289	44.729	1.00 20.16	AAAA
MOTA	540	CD1	ILE	A 3	66		39.528	23.523	45.228	1.00 19.68	AAAA
ATOM	541	С	ILE	E A	66		40.558	23.261	41.023	1.00 22.68	AAAA
MOTA	542	õ	ILE		66		40.636	24.425	40.665	1.00 23.19	AAAA
	543	N	ASN		67		41.036	22.262	40.295	1.00 22.96	AAAA
MOTA											
MOTA	544	CA	ASN		67		41.698	22.545	39.029	1.00 23.92	AAAA
ATOM	545	CB	ASN		67		42.292	21.261	38.395	1.00 24.24	AAAA
MOTA	546	CG	ASN	ΙA	67		43.344	20.588	39.289	1.00 23.38	AAAA
MOTA	547	OD1	ASN	I A	67		44.196	21.256	39.859	1.00 23.47	AAAA
MOTA	548	ND2	ASN	I A	67		43.290	19.258	39.392	1.00 23.20	AAAA
ATOM	549	С	ASN		67		40.717	23.216	38.063	1.00 23.82	AAAA
	550	ō	ASN		67		41.123	23.996	37.204	1.00 24.63	AAAA
MOTA										1.00 24.08	
ATOM	551	N	THR		68		39.427	22.928	38.213		AAAA
MOTA	552	CA	THE		68		38.428	23.534	37.343	1.00 25.28	AAAA
MOTA	553	CB	THR	A	68		37.030	22.904	37.525	1.00 24.55	AAAA
ATOM	554	OG1	THR	A	68		37.090	21.500	37.258	1.00 24.64	AAAA
ATOM	555	CG2	THR	. A	68		36.049	23.534	36.564	1.00 23.58	AAAA
ATOM	556	C	THR		68		38.322	25.023	37.664	1.00 26.31	AAAA
ATOM	557	ŏ	THR		68		38.114	25.854	36.771	1.00 26.69	AAAA
		N	LEU		69		38.462	25.351	38.945	1.00 26.59	AAAA
MOTA	558										
MOTA	559	CA	LEU		69		38.381	26.729	39.378	1.00 27.05	AAAA
ATOM	560	CB	LEU		69		38.321	26.807	40.904	1.00 27.15	AAAA
MOTA	561	CG	LEU	Α	69		37.003	26.397	41.551	1.00 25.68	AAAA
ATOM	562	.CD1	LEU	Α	69		37.088	26.491	43.062	1.00 26.30	AAAA
ATOM	563	CD2	LEU	A	· 69		35.933	27.316	41.044	1.00 26.14	AAAA
ATOM	564	С	LEU		69		39.570	27.508	38.867	1.00 28.44	AAAA
ATOM	565	ō	LEU		69		39.425	28.619	38.356	1.00 28.59	AAAA
	566	N	MET		70		40.748	26.914	39.009	1.00 29.31	AAAA
MOTA					70		41.981	27.536	38.571	1.00 29.89	AAAA
MOTA	567	CA	MET								
ATOM	568	CB	MET		70		43.160	26.692	39.044	1.00 31.04	AAAA
MOTA	569	CG	MET		70		43.164	26.528	40.562	1.00 31.79	AAAA
MOTA	570	SD	MET	Α	70		44.608	25.684	41.183	1.00 32.58	AAAA
MOTA	571	CE	MET	A	70		45.859	26.820	40.670	1.00 30.82	AAAA
ATOM	572	С	MET	A	70		42.017	27.723	37.057	1.00 30.36	AAAA
ATOM	573	Ō	MET		70		42.462	28.769	36.559	1.00 30.18	AAAA
		- N	GLU		71		41.538	26.719	36.328	1.00 30.34	AAAA
MOTA					71		41.519	26.795	34.874	1.00 30.73	AAAA
ATOM	575	CA	GLU								
ATOM	576	CB	GLU		71		41.140	25.442	34.266	1.00 33.47	AAAA
ATOM	577	CG	GLU		71		41.122	25.430	32.731	1.00 37.11	AAAA
ATOM	578	CD	GLU	A	71		42.513	25.676	32.093	1.00 40.49	AAAA
MOTA	579	OE1	GLU	Α	71		42.570	25.798	30.844	1.00 41.95	AAAA
ATOM	580	OE2	GLU	А	71		43.541	25.738	32.825	1.00 40.74	AAAA
	581	C	GLU		71		40.537	27.851	34.392	1.00 29.78	AAAA
MOTA	582	ō	GLU		71		40.852	28.642	33.508	1.00 27.82	AAAA
MOTA									34.992	1.00 27.02	
MOTA	583	N	ALA		72		39.352	27.855			AAAA
MOTA	584	CA	ALA		72		38.296	28.790	34.635	1.00 29.88	AAAA
MOTA	585	CB	ALA	Α	72		37.022	28.432	35.374	1.00 29.07	AAAA
ATOM	586	С	ALA	А	72		38.667	30.238	34.907	1.00 30.78	AAAA
MOTA	587	0	ALA		72		38.359	31.122	34.108	1.00 31.27	AAAA
	588	N	GLU		73		39.336	30.491	36.023	1.00 31.07	AAAA
MOTA			GLU		73		39.710	31.856	36.346	1.00 31.65	AAAA
MOTA	589								37.785	1.00 30.52	
ATOM	590	CB	GLU		73		40.243	31.954			AAAA
MOTA	591	CG	GLU		73		40.643	33.370	38.198	1.00 28.73	AAAA
ATOM	592	CD	GLU	A	73		41.076	33.484	39.651	1.00 28.77	AAAA
ATOM	593	OE1	GLU	Α	73		40.239	33.260	40.546	1.00 28.94	AAAA
ATOM	594	OE2			73		42.258	33.795	39.906	1.00 28.57	AAAA
									-		

		•		~~			22 461	35.378	1.00 3	3 E 4.	AAAA
ATOM	595	C	GLU A	73		0.726	32.461				
ATOM	596	Q	GLU A	73	4	0.456	33.499	34.767	1.00 3		AAAA
ATOM	597	N	ARG A	74	4.	1.885	31.832	35.214	1.00 3	34.35	AAAA
ATOM	598	CA	ARG A	74	4	2.890	32.428	34.334	1.00	36.04	AAAA
	599	CB	ARG A	74		4.238	31.710	34.482	1.00 3	36 92	AAAA
ATOM							30.313	33.923	1.00		AAAA
ATOM-	600	CG	ARG A	74		4.327					
ATOM	601	CD	ARG A	74	4:	5.508	29.589	34.543	1.00 3		, AAAA
ATOM	602	NE	ARG A	74	4.	5.893	28.404	33.785	1.00 4	42.02	AAAA
ATOM	603	CZ	ARG A	74	4	6.632	28.436	32.675	1.00 4	42.69	AAAA
	604		-ARG A	74		7.071	29.593	32.191	1.00 4		AAAA
ATOM							27.309	32.046	1.00		AAAA
MOTA	605		ARG A	74		6.933					
MOTA	606	С	ARG A	74		2.476	32.532	32.864	1.00		AAAA
ATOM	607	0	ARG A	74	- 4:	2.842	33.493	32.18 7	1.00		AAAA
ATOM	608	N	SER A	75	4:	1.711	31.567	32.367	1.00	36.60	AAAA
ATOM	609	CA	SER A	75	4	1.248	31.622	30.987	1.00	36.82	AAAA
		CB	SER A	75		0.916	30.218	30.478	1.00		- AAAA
ATOM	610						29.723	31.083	1.00		AAAA
ATOM	611	OG	SER A	75		9.736					
ATOM	612	C	SER A	75		9.980	32.476	31.001	1.00		AAAA
ATOM	613	0	SER A	75	3 9	9.401	32.791	29.963	1.00	36.25	AAAA
MOTA	614	N	GLN A	76	3 9	9.568	32.845	32.208	1.00	37.62	AAAA
ATOM	615	CA	GLN A	76	3 1	8.368	33.639	32.427	1.00	37.92	AAAA
	616	CB	GLN A	76		8.613	35.100	32.049	1.00		AAAA
MOTA								32.717	1.00		AAAA
ATOM	617	CG	GLN A	76		7.630	36.048				
MOTA	618	CD	GLN A	76		7.929	36.298	34.199	1.00		AAAA
ATOM	619	OE1	GLN A	76	31	8.226	35.379	34.973	1.00		AAAA
ATOM	620	NE2	GLN A	76	3,	7.833	37.556	34.597	1.00	42.32	AAAA
ATOM	621	С	ĠŁN A	76	3'	7.223	33.064	31.600	1.00	37.75	AAAA
ATOM	622	0	GLN A	76	. 31	6.521	33.789	30.901	1.00	38.13	AAAA
	623	N	SER A	77		7.045	31.749	31.685	1.00	37.52	AAAA
ATOM	624	CA	SER A	77		5.990	31.061	30.950	1.00		AAAA
MOTA						6.537	30.440	29.664	1.00		· AAAA
ATOM	625	CB	SER A	77	_		31.441	28.724	1.00		AAAA
MOTA	6 26	OG	SER A	77		6.851					
MOTA	627	С	SER A	77		5.338	29.960	31.757	1.00		AAAA
ATOM	628	0	SER A	77		5.790	29.620	32.846	1.00		AAAA
ATOM	629	N	VAL A	78	34	4.264	29.412	31.198	1.00		AAAA
ATOM	630	CA	VAL A	78	3.	3.538	28.309	31.812	1.00		AAAA
ATOM	631	CB	VAL A	78	3:	2.027	28.514	31.715	1.00	37.19	AAAA
ATOM	632	CG1	VAL A	78	3:	1.310	27.439	32.497	1.00	36.84	AAAA
ATOM	633		VAL A	78		1.662	29.906	32.201	1.00	37.60	AAAA
	634	c	VAL A	78		3.918	27.089	30.976	1.00	38.28	AAAA
ATOM .				78		3.497	26.959	29.819	1.00		AAAA
ATOM	635	0	VAL A				26.187	31.537	1.00		AAAA
ATOM	636	N	PRO A	79		4.734			1.00		AAAA
ATOM	637	CD	PRO A	79		5.347	26.167	32.869			
ATOM	638	CA	PRO A	79		5.146	24.998	30.797	1.00		AAAA
ATOM	639	CB	PRO A	79	3 (5.127	24.325	31.759	1.00		AAAA
ATOM	640	CG	PRO A	79	3 6	5.655	25.489	32.557	1.00	37.65	AAAA
ATOM	641	C	PRO A	79	33	3.980	24.089	30.434	1.00	37.20	AAAA
	642	ō	PRO A	79		2.958	24.050	31.120	1.00	36.43	AAAA
ATOM	643	N	LYS A	80	_	1.154	23.363	29.338	1.00		AAAA
ATOM							22.423	28.855		37.35	
ATOM	644	CA	LYS A	80		3.160			1.00		AAAA
ATOM	645	CB	LYS A	80		3.757	21.586	27.725			
ATOM	646	CG	LYS A	80		2.928	20.379	27.280	1.00		AAAA
ATOM	647	CD	LYS A	80	3 1	L.835	20.710	26.286	1.00		AAAA
ATOM	.648	CE	LYS A	80	31	1.320	19.402	25.688	1.00		AAAA
ATOM	649	NZ	LYS A	80	3 (.498	19.543	24.450	1.00	40.48	AAAA
	650	c	LYS A	80		2.752	21.515	30.003	1.00	36.85	AAAA
ATOM		ò	LYS A	80		3.610	20.942	30.676	1.00		AAAA
ATOM	651					L.443	21.408	30.217	1.00		AAAA
ATOM	652	N	GLY A	81			20.570	31.268	1.00		AAAA
ATOM	653	CA	GLY A	81		0.903					
ATOM	654	С	GLY A	81		L.110	21.054	32.695	1.00		AAAA
TOM	655	0	GLY A	81		749	20.355	33.644	1.00		AAAA
ATOM	656	N	ALA A	82	31	L.677	22.241	32.867	1.00		AAAA
ATOM	657	CA	ALA A	82		.919	22.743	34.213	1.00	35.02	AAAA
	658	CB	ALA A	82		.076	23.743	34.208	1.00		AAAA
ATOM	659	C .	ALA A	82		.674	23.378	34.797	1.00		AAAA
ATOM				82).451	23.332	36.001	1.00		AAAA
ATOM	660	0	ALA A	- A-	31	,		-			•
			A ***								

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ATOM	661	N	ARG	Δ	83	29.858	23.960	33.932	1.00 34.77	AAAA
ATOM	662		ARG		83	28.637			1.00 35.34	
	663	CB	ARG		83	27.899				AAAA
MOTA									1.00 36.26	AAAA
MOTA	664	CG	ARG		83	27.045			1.00 37.09	AAAA
ATOM	665	CD	ARG		83	26.209			1.00 37.48	AAAA
ATOM	666	NE	ARG		83	25.475			1.00 37.35	AAAA
MOTA	667	CZ	ARG	Α	83	24.711	27.311	36.218	1.00 37.77	AAAA
MOTA	668	NH1	ARG	A	83	24.606	26.204	36.940	1.00 37.29	AAAA
MOTA	669	NH2	ARG	Α.	83	24.040	28.401	36.568	1.00 38.34	AAAA
MOTA	670	С	ARG		83	27.739			1.00 36.30	AAAA
MOTA	671	ō	ARG		83	27.232			1.00 36.17	AAAA
MOTA	672	Ŋ	GLU		84	27.565			1.00 37.19	AAAA
	673	CA	GLU		84	26.721			1.00 37.19	
MOTA						26.466				AAAA.
MOTA	674	CB	GLU		84				1.00 40.55	AAAA
ATOM	675	CG	GLU		84	25.643			1.00 43.12	AAAA
MOTA	676	CD	GLU		84	25.362			1.00 44.98	AAAA
ATOM	677		GLU		84	24.573			1.00 46.36	AAAA
MOTA	678	OE2	GLU	Α	84	25.937	18.532	31.962	1.00 44.94	AAAA
ATOM	679	С	GLU	A	84	27.290	20.657	36.158	1.00 37.07	AAAA
ATOM	680	0	GLU	A	84	26.642	20.555	37.199	1.00 36.17	AAAA
MOTA	681	N	LYS	Α	85	28.506	20.152	35.999	1.00 36,23	AAAA
ATOM	682	CA	LYS		85	29.202			1.00 35.36	AAAA
ATOM	683	CB	LYS		85	30.449			1.00 36.96	AAAA
MOTA	684	CG	LYS		85	31.394			1.00 30.04	AAAA
	685	CD	LYS		85	30.995	16.766		1.00 40.59	
MOTA	686	CE			85		15.719	36.933		AAAA
MOTA			LYS			31.508			1:00 41.88	AAAA
ATOM	687	NZ	LYS		85	32.998	15.817		1.00 42.00	AAAA
MOTA	688	C	LYS		85	29.620	20.202	38.289	1.00 33.86	AAAA
MOTA	689	0	LYS		85	29.576	19.679	39.404	1.00 33.82	AAAA
MOTA	690	N	TYR		86	30.014	21.458	38.097	1.00 32.06	AAAA
MOTA	691	CA	TYR .		86	30.514	22.279	39.194	1.00 29.44	AAAA
MOTA	692	CB	TYR	A	86	31.956	22.683	38.875	1.00 29.97	AAAA
MOTA	693	CG	TYR	A	86	32.872	21.496	38.621	1.00 29.99	AAAA
ATOM	694	CD1	TYR .	A	86	33.281	20:666	39.666	1.00 29.24	AAAA
MOTA .	695	CE1	TYR .	A	86	34.126	19.582	39.437	1.00 29.85	AAAA
MOTA	696	CD2	TYR .	A	86	.33.329	21.204	37.329	1.00 30.16	AAAA
ATOM	697	CE2	TYR .	A	86	34.173	20.118	37.087	1.00 29.61	AAAA
ATOM	698	CZ	TYR		86 '	34.570	19.313	38.148	1.00 29.79	AAAA
ATOM	699	ОН	TYR		86	35.414	18.253	37.923	1.00 29.48	AAAA
MOTA	700	c	TYR		86	29.705	23.509	39.572	1.00 27.81	AAAA
MOTA	701	ō	TYR		86	30.052	24.202	40.524	1.00 27.56	AAAA
	702	N	ASN		87	28.642	23.784	38.828	1.00 26.60	
ATOM										AAAA
ATOM	703	CA	ASN A		87	27.777	24.924	39.111	1.00 26.56	AAAA
ATOM	704	CB	ASN A		87	27.172	24.772	40.508	1.00 26.39	AAAA
MOTA	705	CG	ASN A		87	25.863	25.544	40.684	1.00 26.64	AAAA
MOTA	706		ASN A		87	25.335	25.632	41.790	1.00 26.84	AAAA
ATOM	707	ND2	ASN A		87	25.330	26.084	39.597	1.00 26.33	AAAA
ATOM	708	C	ASN A	A (87	28.587	26.217	39.024	1.00 26.40	AAAA
MOTA	709	0	ASN A	A (87	28.430	27.129	39.832	1.00 24.80	AAAA
ATOM	710	N	ILE A	A 6	88	29.448	.26.273	38.015	1.00 27.57	AAAA
ATOM	711	CA	ILE A		88	30.330	27.409	37.767	1.00 27.88	AAAA
ATOM	712	CB	ILE A		88	31.817	26.932	37.648	1.00 27.38	AAAA
ATOM	713		ILE A		88	32.684	27.994	36.986	1.00 26.34	AAAA
	714		ILE A		88	32.354	26.543	39.026	1.00 28.35	AAAA
MOTA										
MOTA	715		ILE A		88	32.356	27.671	40.042	1.00 27.78	AAAA
MOTA	716		ILE A		88	29.946	28.110	36.472	1.00 29.17	AAAA
ATOM	717		ILE A		88	29.530	27.469	35.515	1.00 29.75	AAAA
MOTA	718		GLY A		39	30.092	29.429	36.443	1.00 29.96	AAAA
ATOM	719		GLY Y		39	29.791	30.162	35.229	1.00 30.24	AAAA
ATOM	720	C	GLY A	٤ ،	39.	28.430	30.805	35.242	1.00 30.44	AAAA
ATOM	721		GLY A	. 8	39	28.177	31.769	34.514	1.00 31.14	AAAA
MOTA	722		GLY A		90	27.542	30.268	36.061	1.00 30.00	AAAA .
ATOM	723		GLY A		90	26.221	30.841	36.129	1.00 30.52	AAAA
atom Atom	724		GLY A		90	26.283	32.262	36.661	1.00 31.09	AAAA
	725		GLY A		90	27.356	32.795	36.962	1.00 30.34	AAAA
MOTA								36.768	1.00 31.09	AAAA
atom	726	N :	TYR A	. ,	1	25.112	32.873		1.00 31.03	WWW

n molf	727	CA	TYR	Δ	91	24.977	34.213	37.290	1.00 31.27	AAAA
ATOM		CB	TYR		91	23.515	34.634	37.195	1.00 31.82	AAAA
MOTA	728		_		91	23.169	35.825	38.047	1.00 31.81	AAAA
ATOM	729	CG	TYR			23.536	37.108	37.670	1.00 32.44	AAAA
MOTA	730	CD1			91	23.250	38.203	38.475	1.00 31.88	AAAA
ATOM	731	CE1	TYR		91		35.663	39.254	1.00 32.63	AAAA
ATOM	732	CD2	TYR		91	22.505	36.754	40.068	1.00 32.60	AAAA
MOTA	733	CE2	TYR		91			39.668	1.00 31.59	AAAA
MOTA	734	CZ	TYR		91	22.589	38.016	40.450	1.00 31.94	AAAA
MOTA	735	OH	TYR		91	22.283	39.094		1.00 31.54	AAAA
MOTA	736	С	TYR		91	25.384	34.202	38.753		AAAA
MOTA	737	0	TYR	Α	91	26.075	35.105	39.233	1.00 31.21	AAAA
ATOM	738	N	GLU	Α	92	24.925	33.158	39.438	1.00 31.51	
MOTA	739	CA	GLU	Α	92	25.143	32.941	40.865	1.00 32.70	AAAA
MOTA	740	CB	GLU	Α	92	24.463	31.626	41.268	1.00 33.55	AAAA
ATOM	741	ĊĠ	GLU	A	92	24.174	31.495	42.747	1.00 34.16	AAAA
ATOM	742	CD	GLU	A	92	23.311	30.278	43.087	1.00 35.31	AAAA
ATOM	743	OE1	GLU	Α	92 .	23.857	29.148	43.152	1.00 34.30	AAAA
ATOM	744	OE2	GLU	Α	92	22.076	30.466	43.275	1.00 35.36	AAAA
MOTA	745	C	GLU		92	26.619	32.902	41.248	1.00 33.02	AAAA
ATOM	746	0	GLU		92	27.073	33.623	42.140	1.00 32.91	AAAA
MOTA	747	N	ASN		93	27.358	32.049	40.550	1.00 32.84	AAAA
ATOM	748	CA	ASN		93	28.785	31.861	40.777	1.00 31.92	AAAA
ATOM	749	СВ	ASN		93	29.015	30.437	41.278	1.00 31.18	AAAA
MOTA	750	CG	ASN		93	27.948	29.994	42.259	1.00 30.34	AAAA
	751		ASN		93	27.723	30.642	43.271	1.00 31.20	AAAA
ATOM	752		ASN		93	27.284	28.892	41.955	1.00 29.02	AAAA
ATOM	753	C	ASN		93	29.442	32.052	39.411	1.00 30.84	AAAA
ATOM	754	ō	ASN		93	29.823	31.082	38.758	1.00 30.82	AAAA
ATOM	755	N	PRO		94	29.605	33.309	38.975	1.00 29.56	AAAA
ATOM	756	CD	PRO		94	29.312	34.590	39.626	1.00 29.03	AAAA
MOTA	757	CA	PRO		94	30.209	33.564	37.671	1.00 28.89	AAAA
ATOM	758	CB	PRO		94	29.890	35.045	37.416	1.00 28.22	AAAA
ATOM	759	CG	PRO		94	28.839	35.377	38.435	1.00 29.50	AAAA
ATOM	760	C	PRO		94 .	31.698	33.351	37.664	1.00 28.25	AAAA
ATOM .	761	ò	PRO		94	32.308	32.996	38.671	1.00 28.21	AAAA
MOTA	762	N	VAL		95	32.257	33.593	36.488	1.00 27.36	AAAA
MOTA	763	CA	VAL		95	33.676	33.530	36.247	1.00 26.24	AAAA
MOTA	764	CB	VAL		95	33.945	33.289	34.741	1.00 26.10	AAAA
MOTA	765		VAL		95	35.373	33.717	34.357	1.00 25.47	AAAA
MOTA	766	CG2	VAL		95	33.736	31.826	34.434	1.00 25.59	AAAA
ATOM	767	C	VAL		95	34.178	34.919	36.647	1.00 26.56	AAAA
ATOM	768	ō	VAL		95	33.560	35.937	36.307	1.00 27.18	AAAA
ATOM	769	N	SER		96	35.280	34.966	37.382	1.00 25.23	AAAA
MOTA	770	CA	SER		96	35.858	36.237	37.790	1.00 24.51	AAAA
ATOM	771	CB	SER		96	34.935	36.961	38.774	1.00 23.22	·AAAA
ATOM	772	OG	SER		96	34.941	36.297	40.014	1.00 19.76	AAAA
ATOM	773	C	SER		96	37.169	35.920	38.485	1.00 24.84	AAAA
ATOM	774	ō	SER		96	37.590	34.764	38.530	1.00 25.97	AAAA
ATOM	775	N	TYR		97	37.824	36.933	39.030	1.00 24.02	AAAA
ATOM			TYR		97	39.047	36.664	39.744	1.00 24.55	AAAA
MOTA	776	CA	TYR		97	40.071	37.762	39.504	1.00 23.94	AAAA
MOTA	777	CG	TYR		97	40.682	37.636	38.128	1.00 23.72	AAAA
ATOM	778		TYR		97	40.177	38.341	37.039	1.00 23.11	AAAA
ATOM	779				97	40.700	38.136	35.758	1.00 23.50	AAAA
ATOM	780		TYR		97	41.717	36.735	37.903	1.00 22.25	AAAA
ATOM	781	CD2	TYR		97	42.236	36.526	36.640	1.00 22.86	AAAA
ATOM	782	CE2	TYR		97	41.730	37.217	35.572	1.00 23.56	AAAA
ATOM	783	CZ	TYR			42.232	36.941	34.318	1.00 24.06	AAAA
ATOM	784	OH	TYR		97 97	38.800	36.436	41.228	1.00 25.08	AAAA
MOTA	785	Ċ	TYR		97 97		36.266	42.009	1.00 26.91	AAAA
MOTA	786	0	TYR		97	39.739	36.406	41.589	1.00 24.73	AAAA
ATOM	787	N	ALA		98	37.522	36.159	42.951	1.00 24.50	AAAA
ATOM	788	CA	λLA		98	37.083	36.925	43.235	1.00 24.48	AAAA
MOTA	789	CЗ	ALA		98	35.800	34.661	43.088	1.00 23.95	AAAA
MOTA	790	C	ALA		98	36.824		44.171	1.00 24.21	AAAA
MOTA	791	0 .	ALA		98	36.929	34.100	41.976	1.00 23.10	AAAA
ATOM	792	N	MET	Α	99	36.502	34.011	4270	1.00 20.10	
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Figure 19-13

ATOM	793	CA	MET A	99		36.208	32.584	42.000	1.00	22.61	AAAA
MOTA	794	CB	MET A	99		35.855	32.089	40.597	1.00	23.25	· AAAA
ATOM	795	CG		99		37.009	32.063	39.607		23.22	AAAA
ATOM	796	SD	MET A	99		36.360	31.808			25.21	AAAA
ATOM	797	CE	MET A	99		35.328	30.374			22.04	AAAA
ATOM	798	c	MET A	99		37.319	31.720			21.80	AAAA
		õ	MET A	99		37.052	30.695				
MOTA	799		-	_						21.29	AAAA
MOTA	800	N				38.567	32.111			21.87	AAAA
MOTA	801	CA				39.650	31.322			21.11	AAAA
ATOM	802	CB	PHE A			40.388	30.552			20.25	AAAA
MOTA	803	CG			-	41.451	29.648			20.14	AAAA
MOTA	804		1 PHE A			41.114	28.462			20.49	AAAA
MOTA	805		2 PHE A			42.785	30.050			19.82	AAAA
MOTA	806	CE:	l PHE A	100		42.090	27.695	43.646	1.00	19.54	AAAA
ATOM	807	CE	PHE A	100		43.755	29.300	43.001	1.00	19.22	AAAA
ATOM	808	CZ	PHE A	100		43.410	28.122	43.641	1.00	19.47	AAAA
MOTA	809	С	PHE A	100		40.649	32.161	43.743	1.00	21.37	AAAA
MOTA	810	0	PHE A	100	•	40.959	31.822	44.887	1.00	21.26	AAAA
ATOM	811	N	THR A			. 41.142	33.252	43.161	1.00	20.94	AAAA
ATOM	812	CA	THR A			42.119	34.097			21.95	AAAA
ATOM	813	CB	THR A			42.691	35.181	42.905		22.21	AAAA
ATOM	814		THR A			43.511	34.552	41.917		22.90	AAAA
ATOM	815		THR A			43.535	36.186	43.667		21.38	AAAA
ATOM	816	C	THRA			41.584	34.755	45.117		22.60	
	817	ò	THR A			42.248	34.723	46.147	1.00		AAAA
ATOM		N	GLY A			40.394	35.343	45.049	1.00		
ATOM	818 819		GLY A			39.826	35.972	46.227	1.00		AAAA
MOTA		CA								22.03	AAAA
ATOM	820	C	GLY A			39.340	34.928	47.221			AAAA
MOTA	821	0	GLY A	-		39.433	35.104	48.439		20.02	AAAA
MOTA	822	N	SER A			38.816	33.833	46.677		21.86	AAAA
ATOM	823	CA	SER A			38.311	32.719	47.466	1.00		AAAA
ATOM	824	CB	SER A			37.699	31.668	46.557	1.00	_	AAAA
ATOM	825	OG	SER A			36.604	32.216	45.857	1.00		AAAA
MOTA	826	, C	SER A			39.450	32.098	48.229	1.00		AAAA
MOTA	827	0	SER A			39.314	31.806	49.412	1.00		AAAA
MOTA	828	N	SER A			40.578	31.898	47.545	1.00		AAAA
ATOM	829	CA	SER A			41.746	31.305	48.183	1.00		A A AA
ATOM	830	CB	SER A			42.862	31.070	47.172	1.00		AAAA
MOTA	831	OG	SER A			42.441	30.169	46.175	1.00		AAAA
MOTA	832	С	SER A	104		42.254	32.230	49.256	1.00		AAAA
ATOM	833	0	SER A			42.707	31.794	50.307	1.00		AAAA
MOTA	834	N	LEU A	105		42.160	33.518	48.970	1.00		AAAA
MOTA	835	CA	LEU A	105		42.626	34.541	49.870	1.00	21.70	AAAA
MOTA	836	CB	LEU A	105		42.524	35.882	49.159	1.00	21.89	AAAA
MOTA	837	CG	LEU A	105	٠	43.332	37.038	49.718	1.00	23.64	AAAA
ATOM	838	CD1	LEU A	105		44.830	36.692	49.639	1.00	22.01	AAAA
ATOM	839	CD2	LEU A	105		43.004	38.304	. 48.919	1.00 2	23.60	AAAA
ATOM	840	С	LEU A	105		41.767	34.525	51.131	1.00 2	22.29	AAAA
ATOM	841	0	LEU A	105		42.277	34.595	52.249	1.00 2	21.95	AAAA
ATOM	842	N	ALA A	106		40.458	34.429	50.934	1.00 2	22.23	AAAA
ATOM	843	CA	ALA A				34.394	52.042	1.00 2	22.32	AAAA
MOTA	844	CB	ALA A I			38.068	34.472	51.526	1.00 2		AAAA
MOTA	845	C	ALA A				33.126	52.840	1.00 2		AAAA
ATOM	846	ō	ALA A 1			39.578	33.145	54.061	1.00 2		AAAA
ATOM	847	N	THR A 1			40.011	32.032	52.144	1.00 2		AAAA
MOTA	848	CA	THR A 1			40.209	30.732	52.779	1.00 2		AAAA
	849	CB	THR A 1			40.170	29.571	51.749	1.00		AAAA
ATOM	850	CG1				38.903	29.553	51.083	1.00		
MOTA						40.360	28.242	52.455	1.00		AAAA
MOTA	851		THR A 1					53.561			AAAA
ATOM	852	C	THR A 1			41.516	30.630		1.00 2		AAAA
ATOM	853	0	THR A 1			41.537	30.040	54.646	1.00 2	_	AAAA
ATOM	854	N	GLY A 1			42.601	31.176	53.003	1.00 2	_	AAAA
MOTA	855	CA	GLY A 1			43.878	31.145	53,684	1.00 1		AAAA
ATOM ·	856	С	GLY A 1			43.739	31.933	54.972	1.00 1		AAAA
ATOM	857	0	GLY A 1			44.335	31.600	55.998	1.00 1		AAAA
ATOM	858	N	SER A 1	.09		42.909	32.969	54.929	1.00 1	.v.56	, AAAA

MOTA	859	CA	SER	Α	109		42.683	33.805	56.098	1.00		AAAA
	860	CB	SER				41.899	35.058	55.707	1.00	20.27	AAAA
ATOM .		OG	SER				42.618	35.803	54.746	1.00	21.80	AAAA
MOTA	861						41.955	33.066	57.219	1.00	19.61	AAAA
MOTA	862	C	SER						58.388	1.00		AAAA
ATOM	863	0	SER				42.078	33.426		1.00		AAAA
ATOM -	864	N	THR	Α	110		41.186	32.042	56.866			
MOTA	865	CA	THR	Α	110		40.493	31.288	57.891	1.00		AAAA
ATOM	866	CB	THR				39.365	30.438	57.304	1.00	20.62	AAAA
			THR				38.236	31.284	57.050	1.00	20.80	AAAA
MOTA	867						38.974	29.313	58.262	1.00	20.53	AAAA
ATOM	868		THR						58.601	1.00		AAAA
MOTA	869	С			110		41.504	30.420		1.00		AAAA
ATOM	870	0	THR	Α	.110		41.455	30.268	59.822			
MOTA	871	N	VAL	Α	111	-	42.431	29.855	57.832 ⁻			AAAA
	872	CA	VAL				43.480	29.053	58.423	1.00	21.03	AAAA
ATOM	873	CB	VAL				44.318	28.323	57.345	1.00	21.05	AAAA
MOTA							45.537	27.644	57.983	1.00	19.91	-AAAA
MOTA	874		VAL					27.281	56.648	1.00		AAAA
MOTA	875		VAL				43.460			1.00		AAAA
ATOM	876	С	VAL	Α	111		44.374	30.005	59.232			
ATOM	877	0	VAL	Α	111		44.825	29.671	60.331	1.00		AAAA
	878	N			112		44.612	31.204	58.712	. 1.00	21.62	AAAA
ATOM	879	CA	GLN				45.449	32.133	59.452	1.00	21.89	AAAA
MOTA							45.630	33.450	58.690	1.00	22.50	AAAA
MOTA	880	CB	GLN						57.335	1.00		AAAA
ATOM	881	CG	GLN	A	112		46.288	33.283				AAAA
ATOM	882	CD	GLN	A	112		46.414	34.578	56.569		23.18	
ATOM	883	OE1	GLN	Α	112		47.389	35.310	56.722		23.86	AAAA
	884		GLN				45.413	34.879	55.752		21.90	AAAA
ATOM	885	C			112		44.766	32.383	60.774	1.00	21.84	AAAA
ATOM							45.389	32.316	61.835	1.00	22.47	AAAA
ATOM	886	0			112			32.651	60.700		21.34	AAAA
ATOM	887	N			113		43.468		61.884		20.84	AAAA
MOTA	888	CA			113	-	42.682	32.934				AAAA
ATOM	889	CB	ALA	Α	113		41.244	33.172	61.504		18.52	
ATOM	890	С	ALA	А	113		42.795	31.782	62.865		21.75	AAAA
	891	ō			113		42.880	31.985	64.084	1.00	22.24	AAAA
ATOM					114		42.797	30.569	62.329	1.00	22.54	AAAA
ATOM	892	N						29.393	63.160	1.00	23.16	AAAA
MOTA	893		ILE				42.891		62.352		23.33	AAAA
ATOM	894	CB			114		42.557	28.146				AAAA
ATOM	895	CG2	ILE	Α	114		42.939	26.912	63.106		23.80	
MOTA	896	CG1	ILE	Α	114		41.058	28.130	62.047		23.48	AAAA
	897		ILE				40.610	26.951	61.204		22.08	AAAA
MOTA	898	c			114		44.268	29.270	63.792	1.00	24.33	AAAA
ATOM					114		44.373	29.013	64.990	1.00	25.30	AAAA
ATOM	899	0					45.319	29.490	63.002	1.00	24.96	AAAA
MOTA	900	N			115			29.395	63.503		26.61	AAAA
MOTA	901	CA			115		46.699				24.75	AAAA
ATOM	902	CB	GLU	Α	115		47.708	29.753	62.406			
MOTA	903	CG	GLU	Α	115		47.444	29.033	61.103		25.80	AAAA
	904	CD	GLU	А	115		48.471	29.323	60.030	1.00	26.07	AAAA
MOTA	905		GLU				48.911	30.484	59.940	1.00	27.15	AAAA
ATOM			GLU				48.819	28.402	59.260	1.00	25.45	AAAA
ATOM	906							30.340	64.680	1.00	27.89	AAAA
MOTA	907	С			115		46.877				28.04	AAAA
ATOM	908	0	GLU	Α	115		47.480	29.975	65.695			AAAA
ATOM	909	N	GLU	Α	116		46.337	31.552	64.531	1.00	29.15	
	910	CA	GLU	Α	116		46.408	32.579	65.563	1.00	29.42	AAAA
MOTA	911	CB			116		45.751	33.871	65.082		28.26	AAAA
ATOM							46.482	34.529	63.945	1.00	28.93	AAAA
MOTA	912	CG			116	•		34.937	64.318	1.00	28.32	AAAA
MOTA	913	CD	GLU	A	116		47.902		65.123	1 00	27.68	AAAA
ATOM	914	OE1	GLU	Α	116		48.081	35.878				
ATOM	91,5	OE2	GLŲ	Α	116		48.838	34.297	63.810		27.38	AAAA
	916	C			116		45.737	32.126	66.845		29.77	AAAA
ATOM	917	ō			116		46.338	32.196	67.920		30.29	AAAA
ATOM							44.492	31.665	66.72 7	1.00	29.64	· AAAA
ATCM	918	Ŋ			117			31.204	67.887		29.33	AAAA
ATOM	919	CA			117		43.741			1 00	28.89	AAAA
ATOM	920	CB			117		42.425	30.552	67.480			
	921	CG	PHE	Α	117		41.604	30.087	68.651		28.93	AAAA
MOTA	922		PHE	Α	117		41.010	31.010	69.510		28.42	AAAA
ATOM		CD3	Dur	7	117		41.441	28.723	68.910		29.06	AAAA
atom	923	CD2	LUD	7	117		40.261	30.588	70.610	1.00	28.68	AAAA
ATOM	924	CEL	PHE	A	117		40.201	50.500	•			•

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» mow	925	CES	. PHE	3	117		40.695	28.284	70.009	1.00 29.16	AAAA
MOTA									70.862	1.00 29.03	AAAA
ATOM	926	CZ			117		40.103	29.227			
MOTA	927	С	PHE	A	117		44.545	30.195	68.671	1.00 29.22	AAAA
ATOM	928	0	PHE	Α	117		44.677	30.315	69.884	1.00 30.29	AAAA
ATOM	929	N	LEU	Α	118		45.066	29.195	67.965	1.00 29.24	AAAA
	930	CA			118		45.864	28.145	68.576	1.00 29.50	AAAA
ATOM								27.047	67.550	1.00 28.57	AAAA
MOTA	931	CB			118		46.182				AAAA
MOTA	932	CG			118		44.962	26.296	66.989	1.00 28.16	
MOTA	933	CD1	LEU	Α	118		45.421	25.090	66.191	1.00 25.58	AAAA
MOTA	934	CD2	LEU	A	118		44.053	25.846	68.128	1.00 27.64	, AAAA
ATOM	935	С	LEU	А	118		47.150	28.649	69.227	1.00 30.14	AAAA
	936	ŏ			118		47.727	27.954	70.056	1.00 29,94	AAAA
MOTA								29.845	68.847	1.00 31.36	AAAA
MOTA	937	N			119		47.602				
ATOM	938	CA			119		48.798	30.451	69.448	1.00 32.52	AAAA
MOTA	939	CB	LYS	Α	119		49.396	31.539	68.559	1.00 32.38	AAAA
ATOM	940	CG	LYS	A	119		49.882	31.108	67.199	1.00 33.03	AAAA
ATOM	941	CD	LVS	Δ	119		50.371	32.321	66.411	1.00 32.74	AAAA
		CE			119		50.681	31.939	64.972	1.00 33.94	AAAA
MOTA	942							33.099	64.152	1.00 34.93	AAAA
MOTA	943	NZ			119		51.125				AAAA
ATOM	944	С	LYS	Α	119		48.385	31.143	70.744	1.00 33.74	
ATOM	945	0	LYS	А	119		49.218	31.748	71.413	1.00 34.85	AAAA
MOTA	946	N	GLY	Α	120		47.096	31.079	71.073	1.00 33.68	AAAA
ATOM	947	CA			120		46.600	31.736	72.263	1.00 33.69	AAAA
	948	C			120		45.987	33.110	71.988	1.00 34.11	AAAA
MOTA							45.588	33.802	72.932	1.00 33.65	AAAA
MOTA	949	0			120				70.717	1.00 33.58	AAAA
ATOM	950	N			121		45.904	33.513			AAAA
MOTA	951	CA			121		45.326	34.820	70.368	1.00 33.35	
MOTA	952	CB	ASN	A	121	•	46.194	35.537	69.341	1.00 33.18	AAAA
ATOM	953	CG	ASN	Α	121		47.570	35.828	69.859	1.00 34.31	AAAA
ATOM	954	OD1	ASN	Α	121		48.333	34.921	70.154	1.00 35.67	AAAA
ATOM	955	ND2	ASN	A	121		47.897	37.096	69.975	1.00 34.18	AAAA
	956	c			121		43.888	34.805	69.839	1.00 32.85	AAAA
MOTA		ō			121		43.304	33.751	69.599	1.00 32.78	AAAA
MOTA	957						43.338	36.003	69.655	1.00 32.47	AAAA
ATOM	958	N	VAL						69.148	1.00 30.89	AAAA
ATOM	959	CA	VAL				41.980	36.200			AAAA
MOTA	960	CB	VAL				41.182	37.145	70.070	1.00 31.05	
ATOM	961	CG1	VAL	Α	122		39.831	37.423	69.489	1.00 30.95	AAAA.
MOTA	962	CG2	VAL	Α	122		41.038	36.516	71.440	1.00 31.19	AAAA
MOTA	963	С	VAL	Α	122		42.056	36.805	67.750	1.00 30.19	AAAA
ATOM .	964	0	VAL	Α	122		42.694	37.840	67.535	1.00 31.28	AAAA
	965	N	ALA				41.405	36.147	66.800	1.00 28.62	AAAA
MOTA		CA	ALA				41.415	36.589	65.421	1.00 26.49	AAAA
ATOM	966						42.323	35.708	64.599	1.00 26.51	AAAA
MOTA	967	CB	ALA					36.570	64.836	1.00 25.59	AAAA
MOTA	968	С	ALA				40.038			1.00 26.27	AAAA
ATOM	969	0	ALA				39.173	35.814	65.252		
ATOM	970	N	PHE	A	124		39.848	37.421	62.847	1.00 25.44	AAAA
MOTA	971	CA	PHE	Α	124		38.590	37.534	62.156	1.00 23.87	AAAA
ATOM	972	CB	PHE	A	124		37.832	38.779	62.646	1.00 23.58	AAAA
ATOM	973		PHE	Α	124		36.591	39.119	62.841	1.00 23.71	AAAA
	974		PHE				35.668	38.140	62.495	1.00 23.44	AAAA
ATOM.		_		_			36.311	40.449	62.498	1.00 23.75	AAAA
MOTA	975	CDZ	PHE	Ţ	124				61.823	1.00 23.31	AAAA
MOTA	976	CEI	PHE	ų	124		34.479	38.483		1.00 21.71	AAAA
MOTA	977	CE2	PHE	A	124		35.131	40.796	61.833		
ATOM ·	978	CZ	PHE	A	124		34.217	39.815	61.497	1.00 22.35	AAAA
ATOM	979	С	PHE				38.95 1	37.673	61.700	1.00 23.26	AAAA
ATOM	980	0	PHE	А	124		39.720	38.555	61.323	1.00 22.29	AAAA
	981	N	ASN				38.427	36.759	60.897	1.00 23.24	АААА
MOTA	982	CA	ASN	2	125		38.622	36.785	59.457	1.00 21.08	AAAA
MOTA			ASN				39.181	35.470	58.951	1.00 19.90	AAAA
MOTA	983	CB	VOIA	Ţ	125				57.454	1.00 20.64	AAAA
ATOM	984	CG	ASN	A	143		39.098	35.360	56.748	1.00 21.63	AAAA
MOTA	985	OD1	ASN	А	125		39.389	36.317			
ATOM	986	ND2	ASN	A	125		38.721	34.190	56.956	1.00 19.93	AAAA
ATOM	987	С	ASN	A	125		37.269	37.059	58.813	1.00 20.19	AAAA
ATOM	988	ō	ASN	А	125		36.469	36.148	58.579	1.00 19.21	AAAA
	989	N	PRO	À	126		36.991	38.340	58.543	1.00 19.14	AAAA
MOTA		CD	PRO				37.893	39.460	58.858	1.00 19.22	AAAA
ATOM	990	<i>-</i>	- 110	^			3	,,,,,,,	•	_	•

ATOM	991	CA	PRO A	126	35.766	38.849	57.932	1.00 19.52	AAAA
MOTA	992	CB	PRO A		36.005	40.359	57.941	1.00 18.55	AAAA
			PRO A		37.511	40.465	57.799	1.00 17.97	AAAA
ATOM	993	CG						1.00 17.37	
ATOM	994	C	PRO A		35.456	38.313	56.526		AAAA
MOTA	. 995	0	PRO A		34.303	38.349	56.080	1.00 19.68	AAAA
ATOM	996	. N	ALA A	127	36.477	37.814	55.835	1.00 18.17	AAAA
ATOM	997	CA	ALA A	127	36.283	37.314	54.481	1.00 17.66	AAAA
ATOM	998	CB	ALA A		37.547	37.520	53.658	1.00 17.08	AAAA
	999	c	ALA A		35.875	35.857	54.443	1.00 17.46	AAAA
MOTA							53.409	1.00 18.92	
ATOM	1000	0	ALA A		35.438	35.359			AAAA
ATOM	1001	N	GLY A		36.019	35.180	55.570	1.00 15.94	AAAA
MOTA	1002	CA	GLY A	128	35.685	33.780	55.642	1.00 15.45	AAAA
ATOM	1003	С	GLY A	128	34.226	33.593	55.955	1.00 16.08	AAAA
ATOM	1004	0	GLY A		33.485	34.557	55.997	1.00 15.43	AAAA
ATOM	1005	N	GLY A		33.821	32.353	56.198	1.00 16.77	AAAA
			GLY A		32.426	32.082	56.462	1.00 17.82	AAAA
MOTA	1006	CA					55.169	1.00 18.64	
ATOM	1007	С	GLY A		31.669	31.822			AAAA
MOTA	1008	0	GLY A	129	30.469	32.051	55.108	1.00 18.48	AAAA
ATOM	1009	N	MET A	130	32.380	31.368	54.137	1.00 20.45	AAAA
ATOM	1010	CA	MET A	130	31.790	31.029	52.826	1.00 21.60	AAAA
ATOM	1011	CB	MET A		32.866	31.117	51.744	1.00 22.02	AAAA
	1012	CG	MET A		33.551	32.472	51.698	1.00 21.75	AAAA
MOTA					34.971	32.567	50.599	1.00 24.75	AAAA
ATOM	1013	SD	MET A						
ATOM	1014	CE	MET A		34.268	32.137	49.048	1.00 24.40	AAAA
MOTA	1015	С	MET A	130	31.328	29.587	53.002	1.00 22.08	AAAA
MOTA	1016	0	MET A	130	31.970	28.641	52.546	1.00 22.98	AAAA
ATOM	1017	N	HIS A	131	30.184	29.452	53.659	1.00 22.25	AAAA
ATOM	1018	CA	HIS A	131	29.618	28.171	54.062	1.00 20.49	AAAA
ATOM	1019	CB	HIS A		28.832	28.421	55.342	1.00 20.00	AAAA
		CG	HIS A		27.679	29.360	55.161	1.00 17.93	AAAA
ATOM	1020			_		29.846	54.043	1.00 17.88	AAAA
ATOM	1021		HIS A		27.091				
ATOM	1022		HIS A		26.952	29.854	56.219	1.00 19.33	AAAA
MOTA	1023	CE1	HIS A	131	25.968	30.607	55.758	1.00 16.99	AAAA
ATOM	1024	NE2	HIS A	131	26.031	30.617	54.441	1.00 17.43	AAAA
ATOM	1025	С	HIS A	131	28.763	27.332	53.141	1.00 19.97	AAAA
ATOM	1026	ō	HIS A		28.330	26.262	53.541	1.00 19.61	AAAA
ATOM	1027	N	HIS A		28.518	27.796	51.923	1.00 20.11	AAAA
					27.673	27.058	50.994	1.00 17.76	AAAA
ATOM	1028	CA	HIS A					1.00 16.76	AAAA
ATOM	1029	CB	HIS A		26.879	28.044	50.127		
ATOM	1030	CG	HIS A	132	25.824	28.815	50.862	1.00 15.35	AAAA
ATOM	1031	CD2	HIS A	132	25.567	30.146	50.920	1.00 14.15	AAAA
ATOM	1032	ND1	HIS A	132	24.804	28.200	51.557	1.00 16.15	AAAA
MOTA	1033	CE1	HIS A	132	23.966	29.119	52.005	1.00 14.13	AAAA
ATOM	1034		HIS A		24.405	30.307	51.632	1.00 14.65	AAAA
	1035	C	HIS A		28.355	26.051	50.065	1.00 17.99	AAAA
MOTA			HIS A		27.742	25.053	49.684	1.00 18.54	AAAA
ATOM	1.36	0						1.00 17.82	AAAA
ATOM	1.537	N	ALA A		29.604	26.305	49.690		
ATOM	1638	CA	ALA A		30.300	25.441	48.742	1.00 18.38	AAAA
ATOM	1039	CB	ALA A	133	31.684	25.961	48.507	1.00 17.53	AAAA
ATOM	1040	С	ALA A	133	30.366	23.970	49.130	1.00 20.92	AAAA
ATOM	1041	0	ALA A	133	30.578	23.633	50.298	1.00 21.79	AAAA
	1042	N	PHE A		30.184	23.086	48.152	1.00 20.58	AAAA
MOTA					30.258	21.663	48.455	1.00 21.38	AAAA
ATOM	1043	CA	PHE A						AAAA
ATOM	1044	CB	PHE A		29.168	20.860	47.731	1.00 19.41	
MOTA	1045	ÇG	PHE A	134	27.772	21.229	48.126	1.00 18.32	AAAA .
ATOM	1046	CD1	PHE A		27.027	22.099	47.357	1.00 19.22	AAAA
ATOM	1047		PHE A		27.193	20.701	49.271	1.00 19.14	AAAA
ATOM	1048		PHE A		25.714	22.438	47.726	1.00 18.56	AAAA
					25.889	21.036	49.644	1.00 17.72	AAAA
ATOM	1049		PHE A				48.866	1.00 18.01	AAAA
MOTA	1050	CZ	PHE A		25.158	21.903			
MOTA	1051	С	PHE A		31.625	21.124	48.081	1.00 22.90	AAAA
ATOM	1052	0	PHE A	134	32.459	21.833	47.544	1.00 23.37	AAAA
ATOM	1053	N	LYS A	135	31.842	19.861	48.390	1.00 24.63	AAAA
ATOM	1054	CA	LYS A		33.095	19.195	48.122	1.00 27.16	AAAA
	1055	CB	LYS A		32.926	17.714	48.480	1.00 28.53	AAAA
MOTA					34.133	16.843	48.292	1.00 31.01	AAAA
ATOM	1056	CG	LYS A	100	24.123	70.047			
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	1057	CD	LYS A	125		33.879	15.472	48.910	1.00 32.75	AAAA
MOTA								50.457	1.00 33.96	AAAA
ATOM	1058	CE	LYS A			33.961	15.495			
MOTA	1059	ΝZ	LYS ?	135		35.371	15.664	50.976	1.00 33.04	AAAA
ATOM	1060	С	LYS A	135		33.577	19.390	46.673	1.00 27.37	AAAA
ATOM	1061	0	LYS A	135		34.769	19.596	46.437	1.00 27.35	AAAA
	1062	N	SER A			32.658	19.354	45.714	1.00 27.32	AAAA
ATOM						33.028	19.527	44.313	1.00 28.31	AAAA
MOTA	1063		SER A							
MOTA	1064	CB	SER A			33.093	18.162	43.626	1.00 28.56	AAAA
ATOM	1065	OG	SER A	136		33.822	17.242	44.417	1.00 29.28	AAAA
MOTA	1066	С	SER A	136		31.993	20.395	43.599	1.00 28.91	AAAA
ATOM	1067	0	SER A	136		31.568	20.080	42.486	1.00 28.78	AAAA
	1068	N	ARG A			31.595	21.502	44.212	1.00 29.08	_AAAA_
ATOM						30.574	22.311	43.576	1.00 29.66	AAAA
MOTA	1069	CA	ARG A					43.657	1.00 31.65	AAAA
MOTA	1070	CB	ARG A			29.259	21.528			
ATOM	1071	CG	ARG A			27.989	22.273	43.355	1.00 33.89	AAAA
ATOM	1072	CD	ARG' A	137		26.862	21.267	43.37 3	1.00 35.93	AAAA
ATOM	1073	NE	ARG A	137		26.961	20.366	42.228	1.00 36.31	AAAA
ATOM	1074	CZ	ARG A	137		26.505	20.660	41.015	1.00 35.99	AAAA
	1075		ARG A			25.915	21.834	40.798	1.00 34.63	AAAA
MOTA						26.650	19.786	40.025	1.00 35.35	AAAA
ATOM	1076		ARG A					44.116	1.00 28.53	AAAA
MOTA	1077	С	ARG A			30.402	23.723			
ATOM	1078	0	ARG A	137		30.418	23.946	45.324	1.00 28.51	AAAA
ATOM	1079	N	ALA A	138		30.247	24.673	43.202	1.00 27.53	AAAA
ATOM	1080	CA	ALA A	138		30.039	26.063	43.581	1.00 27.64	AAAA
ATOM	1081	CB	ALA A	138		30.236	26.984	42.381	1.00 27.87	AAAA
	1082	c	ALA A			28.601	26.130	44.079	1.00 27.27	AAAA
ATOM			ALA A			27.769	25.321	43.671	1.00 28.30	AAAA
MOTA	1083	0					27.080	44.951	1.00 26.16	AAAA
MOTA	1084	N	ASN A			28.292			1.00 25.39	AAAA
MOTA	1085	CA	ASN A			26.945	27.134	45.480		
ATOM	1086	CB	ASN A	139		26.673	25.847	46.282	1.00 24.58	AAAA
MOTA	1087	CG	ASN · A	139		25.343	25.872	47.017	1.00 25.37	AAAA
ATOM	1088	OD1	ASN A	139		24.272	26.017	46.413	1.00 24.20	AAAA
	1089		ASN A			25.408	25.720	48.338	1.00 24.91	AAAA
ATOM			ASN A			26.683	28.358	46.341	1.00 24.90	AAAA
MOTA	1090	č			•	27.346	28.570	47.348	1.00 24.98	AAAA
ATOM	1091	0	ASN A					45.916	1.00 24.46	AAAA
MOTA	1092	N	GLY A			25.702	29.145			AAAA
ATOM	1093	CA	GLY A	140		25.294	30.336	46.625	1.00 22.96	
MOTA	1094	C	GLY A	140		26.383	31.358	46.755	1.00 22.24	AAAA
ATOM	1095	0	GLY A	140		26.663	31.817	47.867	1.00 23.09	AAAA
ATOM	1096	N	PHE A	141		26.992	31.711	45.625	1.00 20.60	AAAA
MOTA	1097	CA	PHE A			28.075	32.700	45.572	1.00 19.43	AAAA
	1098	CB	PHE A			27.758	33.920	46.430	1.00 19.86	AAAA
MOTA	-					26.453	34.577	46.114	1.00 21.18	AAAA
MOTA	1099	CG	PHE A				35.592	46.934	1.00 20.49	AAAA
MOTA	1100		PHE A			25.974			1.00 21.42	
MOTA	1101		PHE A			25.723	34.218	44.985		· AAAA
MOTA	1102	CEl	PHE A	141		24.800	36.242	46.638	1.00 ?2.45	AAAA
ATOM	1103	CE2	PHE A	141		24.540	34.859	44.672	1.00 :1.76	AAAA
MOTA	1104	CZ	PHE A	141		24.072	35.881	45.499	3.05 1.00	AAAA
	1105	Ċ	PHE A			29.396	32.132	46.069	1.00 18.68	AAAA
MOTA	1106	õ	PHE A				32.784	45.944	1.00 19.19	AAAA
ATOM	_					29.367	30.930	46.635	1.00 16.93	AAAA
MOTA	1107	N	CYS A						1.00 16.80	AAAA
MOTA	1108	CA	CYS A			30.594	30.332	47.150	1.00 10.00	
MOTA	1109	CB	CYS A			30.323	29.689	48.509	1.00 16.51	AAAA
ATOM	1110	SG	CYS A	142	• •	29.524	30.826	49.617	1.00 15.01	AAAA
ATOM	1111	С	CYS A			31.227	29.315	46.221	1.00 16.45	AAAA
	1112	ō	CYS A			30.533	28.565	45.556	1.00 15.32	AAAA
MOTA						32.558	29.311	46.190	1.00 18.39	AAAA
MOTA	1113	N	TYR A				28.394	45.362	1.00 18.63	AAAA
MOTA	1114	CA	TYR A			33:340				
ATOM	1115	CB	TYR A			34.298	29.154	44.438	1.00 19.48	AAAA
ATOM	1116	CG	TYR A			33.664	30.214	43.571	1.00 19.55	AAAA
ATOM	1117	CD1	TYR A	143		33.480	31.510	44.043	1.00 20.27	AAAA
	1118		TYR A			32.856	32.473	43.261	1.00 21.63	AAAA
ATOM		CD2	TYR A			33.212	29.910	42.292	1.00 20.14	AAAA
ATOM	1119					32.588	30.863	41.507	1.00 20.82	AAAA
MOTA	1120	_	TYR A				32.135	41.998	1.00 20.90	AAAA
ATOM	1121	CZ	TYR A			32.414			1.00 23.36	AAAA
MOTA	1122	он	TYR A	143		31.787	33.071	41.228	1.00 23.30	AAAA

MOTA	1123	С	TYR	Α	143		34.162	27.490	46.283	1.00		AAAA
ATOM	1124	ō	TYR				34.319	26.289	46.032	1.00		AAAA
ATOM	1125	N	ILE				34.695	28.087	47.344	1.00		AAAA
ATOM	1126	CA	ILE				35.490	27.350	48.315	1.00	19.97	AAAA
MOTA	1127	CB	ILE				36.952	27.861	48.355	1.00		AAAA
ATOM	1128	CG2					37.757	27.088	49.410	1.00	_	AAAA
ATOM	1129		ILE				37.584	27.671	46.965	1.00	20.12	AAAA
ATOM	1130		ILE				39.053	28.072	46.846	1.00	21.05	AAAA
ATOM	1131	c			144		34.833	27.532	49.665	1.00	20.22	AAAA
ATOM	1132		ILE				34.357	28.626	49.981	1.00	19.94	AAAA
ATOM	1133	N	ASN				34.787	26.451	50.440		20.57	AAAA
ATOM	1134	CA	ASN				34.165	26.448	51.770	1.00	20.39	AAAA
MOTA	1135	CB			145	-	33.450	25.114	51.990	1.00	19.39	AAAA
ATOM	1136	CG			145		32.505	25.143	53.171		19.31	AAAA
ATOM	1137		ASN				32.862	25.583	54.263	1.00	21.26	AAAA
ATOM	1138		ASN				31.290	24.667	52.960		17.08	- AAAA
ATOM	1139	С			145		35.236	26.621	52.856		20.17	AAAA
ATOM	1140	ō			145		35.690	25.622	53.421		19.75	AAAA
ATOM	1141	N			146		35.644	27.862	53.148	1.00	20.06	
ATOM	1142	CA			146		36.671	28.075	54.166		20.98	AAAA
MOTA	1143	CB			146		37.019	29.573	54.333		21.78	AAAA
ATOM	1144	CG			146		35.876	30.411	54.882		22.78	AAAA
ATOM	1145		ASN				35.651	30.465	56.091		22.83	AAAA
ATOM	1146		ASN				35.144	31.078	53.983		23.70	AAAA
ATOM		С			146		36.307	27.413	55.496		21.18	AAAA
MOTA	1148	0	ASN	Α	146		37.169	26.823	56.139		21.48	AAAA
ATOM	1149	N			147		35.031	27.476	55.922		20.88	AAAA
ATOM	1150	CD	PRO	Α	147		33.835	28.120	55.358		21.85	AAAA
ATOM	1151	CA	PRO	Α	147		34.674	26.831	57.183		21.42	AAAA
ATOM	1152	CB	PRO	A	147		33.176	27.073	57.261		21.00	AAAA
ATOM	1153	CG	PRO	A	147	-	33.052	28.408	56.605		20.47	AAAA
ATOM	1154	С	PRO.	Α	147		35.015	25.334	57.174		22.79	AAAA
ATOM	1155	0	PRO	A	147		35.650	24.833	58.099		25.69	AAAA
ATOM	1156	N	ALA	A	148		34.603	24:616	56.136		22.34	AAAA
ATOM	1157	CA	ALA	Α	148		34.889	23.193	56.070		22.23	AAAA
ATOM	1158	CB	ALA	Α	148		34.260	22.561	54.825		22.87	AAAA
ATOM	1159	C ·	ALA	A	148		36.378	22.998	56.054		22.33	AAAA AAAA
ATOM	1160	0			148		36.912	22.249	56.861		23.42	AAAA
ATOM	1161	N			149		37.050	23.661	55.122		22.50 21.29	AAAA
ATOM	1162	CA			149		38.505	23.569	55.018		20.46	AAAA
MOTA	1163	CB			149		39.066	24.581	54.002		19.36	AAAA
ATOM	1164		VAL				40.578	24.607	54.085		20.03	AAAA
MOTA	1165		VAL				38.608	24.229	52.593 56.367		21.48	AAAA
MOTA	1166	С			149		39.164	23.848	56.735		22.11	AAAA
ATOM	1167	0			149		40.147	23.197	57.088		21.19	AAAA
MOTA	1168	N	GL.	A	150		38.628	24.826	58.386		21.70	
MOTA	1169	CA			150		39.171	25.176	59.368		22.31	AAAA
MOTA	1170	C .			150		38.973	24.043 23.597	60.026		22.51	AAAA
MOTA	1171	0			150		39.913	23.566	59.453		22.86	AAAA
ATOM	1172	N			151		37.736	22.474	60.346		22.26	AAAA
MOTA	1173	CA	ILE	A	151		37.000	22.124	60.191	1.00	21.51	AAAA
MOTA	1174	CB	ILE	A	151		35.894	20.899	61.019		21.36	AAAA
MOTA	1175	CG2	ILE	A	121		35.542 35.051	23.329	60.627		20.39	AAAA
ATOM	1176		ILE					23.199	60.361		16.88	AAAA
MOTA	1177		ILE				33.576	21.243	60.096		23.29	AAAA
ATOM	1178	С	ILE	A	151		38.265	20.660	61.038		23.88	AAAA
MOTA	1179	0	ILE	A	151		38.786	20.853	58.836		24.13	AAAA
ATOM	1180	N	GLU	A	152		38.435	19.697	58.517		25.01	AAAA
ATOM	1181	CA	GLU	A	152		39.267	19.697	57.010		25.07	AAAA
ATOM	1182	CB	GLU	A	152		39.242	18.886	56.526		24.56	AAAA
MOTA	1183	CG	GLU	A	152		37.910	17.570	57.198		25.00	AAAA
ATOM	1184	CD	GLU	A	152		37.500	17.158	57.011		26.40	AAAA
MOTA	1185	OE1	GLU	A	152		36.345	16.935	57.897		25.00	AAAA
ATOM	1186		GLU	A	152		38,315	19.957	58.965		26.06	AAAA
ATOM	1187	С	GLU	A	152		40.694 41.425	19.035	59.331		26.40	AAAA
MOTA	1188	0	GLU	A	152		41.460	19.000				
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MOTA	1189	N	TYR A	153	41.085	21.225	58.925	1.00 27.30	AAAA
MOTA	1190	CA	TYR A	153	42.422	21.632	59.334	1.00 27.63	AAAA
ATOM	1191	CB	TYR A	153	42.532	23.153	59.268	1.00 26.99	AAAA
MOTA	1192	CG	TYR A	153	43.856	23.719	59.710	1.00 27.03	AAAA
MOTA	1193	CD1	. TYR A	153	44.942	23.790	58.837	1.00 27.78	AAAA
MOTA	1194	CE1	TYR A	153	46.165	24.356	59.250	1.00 28.40	AAAA
ATOM	1195	CD2			44.017	24.215	61.007	1.00 27.52	AAAA
MOTA	1196	CE2	TYR A	153	45.216	24.774	61.425	1.00 27.66	AAAA
MOTA	1197	CZ	TYR A	153	46.284	24.845	60.547	1.00 28.15	AAAA
ATOM	1198	OH	TYR A	153	47.457	25.407	60.974	1.00 28.83	, AAAA
MOTA	1199	С	TYR A		42.618	21.172	60.769	1.00 27.82	AAAA
MOTA	1200	0	TYR A		43.613	20.552	61.110	1.00 27.15	AAAA
MOTA	1201	N	LEU A		41.636	21.487	61.604	1.00 29.25	AAAA.
MOTA	1202	CA	LEU A		41.665	21.138	63.014	1.00 29.35	AAAA
MOTA	1203	CB	LEU A		40.507	21.829	63.715	1.00 30.25	AAAA
MOTA	1204	CG	LEU A		40.685	23.346	63.792	1.00 31.10	AAAA
MOTA	1205		LEU A		39.348	24.020	64.092	1.00 31.24	AAAA
MOTA	1206		LEU A		41.747	23.669	64.852	1.00 29.84	AAAA
MOTA	1207	C	LEU A		41.625	19.639	63.263	1.00 29.73	AAAA
ATOM	1208	0	LEU A		42.313	19.151	64.150	1.00 30.51	AAAA AAAA
MOTA	1209	N	ARG A		40.832	18.903	62.489 62.671	1.00 28.95 1.00 28.94	AAAA
MOTA	1210	CA	ARG A		40.771	17.459 16.820	61.723	1.00 28.94	AAAA
ATOM	1211	CB	ARG A		39.742 38.312	17.312	61.952	1.00 27.82	AAAA
MOTA	1212	CG	ARG A		37.312	16.751	60.955	1.00 27.19	AAAA
MOTA	1213	CD	ARG A		36.804	15.444	61.338	1.00 28.86	AAAA
ATOM	1214 1215	NE CZ	ARG A		35.939	14.742	60.612	1.00 28.93	AAAA
MOTA	1215		ARG A		35.500	15.227	59.459	1.00 29.47	AAAA
MOTA	1217		ARG A		35.486	13.574	61.053	1.00 28.76	AAAA
ATOM ATOM	1218	C	ARG A		42.158	16.853	62.438	1.00 30.20	AAAA
ATOM	1219	ŏ	ARG A		42.572	15.949	63.164	1.00 30.74	AAAA
MOTA	1220	N	LYS A		42.890	17.362	61.447	1.00 30.32	AAAA
MOTA	1221	CA	LYS A		44.224	16.838	61.173	1.00 30.07	AAAA
MOTA	1222	CB	LYS A		44.771	17.373	59.847	1.00 30.26	AAAA
ATOM	1223	CG	LYS A		46.168	16.869	59.525	1.00 30.16	AAAA
ATOM	1224	CD	LYS A	156	46.686	17.368	58.181	1.00 31.19	AAAA
ATOM	1225	CE	LYS A	156	45.884	16.813	56.986	1.00 31.70	AAAA
ATOM	1226	NZ	LYS A	156	45.963	15.324	56.824	1.00 31.20	AAAA
MOTA	1227	С	LYS A	156	45.167	17.202	62.306	1.00 30.08	AAAA
MOTA	1228	0	LYS A	156	46.192	16.550	62.485	1.00 29.16	- AAAA
ATOM	1229	N	LYS A		44.816	18.252	63.053	1.00 30.08	AAAA
ATOM	1230	CA	LYS A		45.608	18.691	64.196	1.00 31.03	AAAA
MOTA	1231	CB	LYS A		45.446	20.201	64.452	1.00 31.81	AAAA
MOTA	1232	CG	LYS A		46.067	21.134	63.419 63.348	1.00 32.12	AAAA
MOTA	1233	CD	LYS A		47.580	21.041	62.226	1.00 31.34 1.00 32.66	AAAA AAAA
MOTA	1234	CE.	LYS A		48.080	21.941 21.921	61.996	1.00 32.00	AAA
ATOM	1235	NZ	LYS A		49.556 45.196	17.923	65.458	1.00 32.74	. `AAA
MOTA	1236	C	LYS A		45.652	18.230	66.558	1.00 31.73	AAAA
ATOM	1237	0.	LYS A GLY A		44.312		65.299	1.00 32.41	AAAA
MOTA	1238	N CA	GLY A		43.901	16.140	66.436	1.00 32.34	AAAA
MOTA	1239	_			42.604	16.429		1.00 32.65	AAAA
MOTA	1240	C	GLY A		42.182	15.604	67.980	1.00 32.85	AAAA
MOTA	1241 1242	N O	PHE A		41.960	17.565	66.932	1.00 33.16	AAAA
ATOM	1242	CA	PHE A		40.712	17.842	67.650	1.00 34.16	AAAA
MOTA	1244	CB	PHE A		40.220	19.281	67.403	1.00 34.81	AAAA
MOTA	1245	CG	PHE A		41.134	20.343	67.965	1.00 34.01	AAAA
ATOM	1245		PHE A		42.327	20.669	67.329	1.00 34.18	AAAA
MOTA	1247		PHE A		40.821	20.981	69.166	1.00 34.61	AAAA
MOTA	1248		PHE A		43.197	21.610	67.874	1.00 33.65	AAAA
MOTA .	1249		PHE A		41.689	21.924	69.718	1.00 34.52	AAAA
ATOM	1250	CZ	PHE A		42.878	22.236	69.065	1.00 33.90	AAAA
ATOM	1251	C	PHE A		39.645	16.840	67.239	1.00 34.04	AAAA
ATOM	1252	ō	PHE A		39.568	16.456	66.068	1.00 34.98	AAAA
ATOM	1253	N	LYS A		38.839	16.403	68.202	1.00 33.36	AAAA
ATOM	1254	CA	LYS A		37.794	15.415	67.936	1.00 33.11	AAAA
A			An				•		•

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				_		20 000	14.140	68.763	1.00 33.97	AAAA
MOTA	1255		LYS			38.060	13.491	68.457	1.00 35.31	AAAA
MOTA	1256	CG	LXS			39.410		69.429	1.00 36.48	AAAA
MOTA	1257	CD	LYS			39.833	12.364		1.00 37.97	AAAA
ATOM	1258	CE	LYS			39.095	11.037	69.243		AAAA
MOTA	1259	NZ	LYS	Α	160	37.636	11.080	69.568	1.00 39.67	AAAA
MOTA	1260	С	LYS	Α	160	36.385	15.941	68.210	1.00 31.68	AAAA
ATOM	1261	0	LYS	A	160	35.405	15.290	67.887	1.00 31.51	
ATOM	1262	N	ARG	Α	161	36.291	17.114	68.819	1.00 31.11	AAAA
ATOM	1263	CA	ARG	Α	161	35.003	17.719	69.114	1.00 30.92	AAAA
MOTA	1264	CB	ARG			34.655	17.592	70.604	1.00 31.78	AAAA
	1265	CG	ARG			34,451	16.157	71.102	1.00 32.91	AAAA
ATOM	1266	CD	ARG			33.994	16.126	72.570	1.00 33.26	AAAA
MOTA		NE	ARG			34.929	16.797	73.476	1.00 34.01	AAAA
MOTA	1267	CZ	ARG			36.183	16.404	73.698	1.00 34.88	AAAA
ATOM	1268		ARG			36.675	15.334	73.081	1.00 34.89	AAAA
MOTA	1269		ARG			36.954	17.084	74.537	1.00 34.71	AAAA
ATOM	1270					35.061	19.185	68.714	1.00 30.28	AAAA
ATOM	1271	С	ARG			35.365	20.059	69.529	1.00 29.86	AAAA
MOTA	1272	0	ARG			_	19.433	67.437	1.00 28.86	AAAA
ATOM	1273	N	ILE			34.774	20.774	66.862	1.00 26.41	AAAA
MOTA	1274	CA	ILE			34.788	20.762	65.464	1.00 26.87	AAAA
ATOM	1275	CB	ILE			35.443		64.872	1.00 26.91	AAAA
MOTA	1276		ILE			35.453	22.160	65.578	1.00 28.19	AAAA
ATOM	1277		ILE			36.877	20.234		1.00 28.24	AAAA
MOTA	1278	CD1	ILE	A	162	37.614	20.090	64.240		AAAA
ATOM	1279	С	ILE	А	162	33.369	21.283	66.731	1.00 24.08	AAAA
MOTA	1280	0	ILE	Α	162	32.485	20.572	66.267	1.00 24.40	AAAA
MOTA	1281	N	LEU	Α	163	33.153	22.519	67.153	1.00 22.25	
ATOM	1282	CA	LEU	Α	163 .	31.838	23.126	67.074	1.00 20.48	AAAA
MOTA	1283	CB	LEU	Α	163	31.408	23.671	68.440	1.00 20.97	AAAA
ATOM	1284	CG	LEU	Α	163	30.099	24.477	68.486	1.00 20.50	AAAA
ATOM	1285	CD1	LEU	A	163	28.998	23.695	67.799	1.00 19.07	AAAA
ATOM	1286		LEU			29.738	24.802	69.950	1.00 19.76	AAAA
ATOM	1287	C	LEU			31.801	24.241	66.055	1.00 18.76	AAAA
	1288	ō			163	32.756	24.986	65.894	1.00 18.41	AAAA
MOTA	1289	Ŋ			164	30.677	24.344	65.368	1.00 17.85	AAAA
MOTA	1290	CA			164	30.496	25.372	64.373	1.00 17.16	AAAA
ATOM	1291	CB			164	30.644	24.768	62.983	1.00 17.45	AAAA
ATOM	1292	CG			164	30.484	25.783	61.900	1.00 17.70	AAAA
MOTA	1292		TYR				26.772	61.701	1.00 16.23	AAAA
MOTA			TYR			31.280	27.734	60.721	1.00 17.35	AAAA
MOTA .	1294		TYR			29.350	25.781	61.092	1.00 17.95	AAAA
MOTA	1295				164	29.173	26.746	60.103	1.00 18.03	AAAA
ATOM	1296	CE2			164	30.138	27.717	59.919	1.00 17.30	AAAA
MOTA	1297	CZ			164	29.955	28.647	58.926	1.00 16.70	AAAA
MOTA	1298	ОН				29.123	26.016	64.514	1.00 15.85	AAAA
MOTA	1299	C			164	28.101	25.351	64.416	1.00 16.44	AAAA
MOTA	1300	0				29.115	27.319	64.743	1.00 15.54	AAAA
MOTA	1301	N			165	27.878	28.088	64.897	1.00 15.71	AAAA
MOTA	1302	CA				27.869	28.819	66.250	1.00 15.18	AAAA
ATOM	1303	CB			165		29.685	66.374	1.00 13.94	AAAA
ATOM	1304	CG2	ILE	A	165	26.621		67.386	1.00 13.94	AAAA
MOTA	1305	CGI	ILE	A	165	28.000	27.797	68.747	1.00 13.94	AAAA
ATOM	1306	CDI	ILE	A	165	28.356	28.421	63.754	1.00 16.00	AAAA
ATOM	1307	С			165	27.808	29.124	63.734	1.00 16.56	AAAA
ATOM	1308	0			165	28.711	29.941		1.00 16.18	AAAA
ATOM	1309	N			166	26.721	29.087	63.001	1.00 16.10	AAAA
ATOM	1310	CA	ASP	Α	166	26.524	29.962	61.865		
ATOM	1311	CB	ASP	Α	166	26.240	29.066	60.651	1.00 18.05	AAAA
ATOM	1312	CG	ASP	А	166	26.238	29.809	59.329	1.00 19.21	AAAA
ATOM	1313	OD1	ASP	A	166	25.353	30.65 <u>9</u>	59.114	1.00 18.36	AAAA
	1314	OD2	ASP	A	166	27.131	29.521	58.495	1.00 19.19	AAAA
MOTA	1315	C			166	25.342	30.904	62.169	1.00 17.57	AAAA
MOTA		ō			166	24.206	30.459	62.321	1.00 17.26	AAAA
ATOM	1316	И.			167	25.605	32.202	62.274	1.00 16.67	AAAA
ATOM	1317	CA	التجرا	Α	167	24.526	33.135	62.562	1.00 16.89	
MOTA	1318		TETT	A A	167	24.923	34.116	63.663	1.00 17.27	AAAA
MOTA	1319	CB			167	25.499	33.529	64.954	1.00 18.37	AAAA
MOTA	1320	CG	∪عبد	A		44.377		. •		
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Figure 19-21

ATOM	1321	CD	1 LEU	A 167		25.760	34.671	65.933	1.00 18.72	AAAA
MOTA	1322			A 167		24.566	32.507	65.547	1.00 17.06	AAAA
	1323	c		A 167		24.146	33.897	61.307	1.00 17.18	AAAA
MOTA										
ATOM	1324	0		A 167		23.390	34.850	61.358	1.00 17.21	AAAA
MOTA	1325	И		A 168		24.683	33.457	60.178	1.00 17.83	AAAA
ATOM	1326	CA	ASP	A 168		24.382	34.067	58.904	1.00 17.84	AAAA
MOTA	1327	CB	ASP	A 168		25.178	33.397	57.807	1.00 20.42	AA , AA
MOTA	1328	CG		A 168	•	25,140	34.162	56.529	1.00 21.41	AAAA
	1329	c		A. 168		22.915	33.783	58.660	1.00 18.35	AAAA
MOTA						22.419	32.722	59.032	1.00 19.62	AAAA
MOTA	1330	0		A 168						
MOTA	1331			A 168	-	26.066	34.972	56.330	1.00 22.42	AAAA
ATOM	1332	OD:	2 ASP	A 168		24.186	33.971	55.746	1.00 21.79	AAAA
MOTA	1333	N	ALA	A 169		22.239	34.717	58.010	1.00 17.98	AAAA
ATOM	1334	CA	ALA	A 169		20.824	34.601	57.708	1.00 17.36	AAAA
ATOM	1335	СВ	ALA	A 169		20.348	35.860	57.007	1.00 17.00	AAAA
ATOM	1336	c		A 169		20.439	33.377	56.887	1.00 18.64	AAAA
	1337	õ		A 169		19.255	33.043	56.819	1.00 19.46	AAAA
ATOM									1.00 18.71	AAAA
MOTA	1338	N		A 170		21.412	32.712	56.262		
MOTA	1339	CA		A 170		21.107	31.518	55.464	1.00 18.43	AAAA
MOTA	1340	С	HIS	A 170		21.802	30.265	55.986	1.00 18.02	AAAA
MOTA	1341	0	HIS	A 170		22.910	30.332	56.514	1.00 17.20	AAAA
ATOM	1342	CB	HIS	A 170		21.539	31.678	54.004	1.00 18.79	AAAA
ATOM	1343	CG		A 170		21.137	32.968	53.386	1.00 17.65	AAAA
	1344			A 170		21.644	34.162	53.828	1.00 18.08	AAAA
MOTA				A 170		21.112	35.081	53.054	1.00 18.95	AAAA
ATOM	1345							52.348		
ATOM	1346			A 170		20.301	33.194		1.00 18.81	AAAA
MOTA	1347			A 170		20.291	34.544	52.140	1.00 19.66	AAAA
MOTA	1348	N	HIS	A 171		21.142	29.124	55.793	1.00 17.53	AAAA
ATOM	1349	CA	HIS	A 171		21.662	27.822	56.193	1.00 16.38	AAAA
ATOM	1350	CB	HIS	A 171		20.644	26.740	55.830	1.00 16.32	AAAA
MOTA	1351	CG	HIS -	A 171		21.157	25.337	55.958	1.00 15.91	AAAA
MOTA	1352			A 171		21.241	24.336	55.051	1.00 14.14	AAAA
	1353			A 171		21.602	24.807	57.151	1.00 16.79	AAAA
MOTA				A 171		21.937	23.543	56.973	1.00 14.91	AAAA
MOTA	1354								1.00 15.45	AAAA
MOTA	1355		HIS.			21.725	23.234	55.709		
MOTA	1356	С		A 171		22.982	27.522	55.509	1.00 16.94	AAAA
ATOM	1357	0		A 171		23.146	27.725	54.318	1.00 18.71	AAAA
ATOM	1358	N	CYS .	A 172		23.926	27.019	56.279	1.00 16.99	AAAA
ATOM	1359	CA	CYS .	A 172		25.237	26.670	55.778	1.00 16.23	AAAA
MOTA	1360	CB	CYS	A 172		26.219	26.721	56.947	1.00 17.89	AAAA
ATOM	1361	SG		A 172		25.638	25.773	58.397	1.00 17.89	AAAA
	1362	c		A 172		25.205	25.271	55.210	1.00 16.57	AAAA
ATOM				A 172		25.947	24.413	55.670	1.00 17.66	AAAA
MOTA	1363	0						54.214	1.00 18.25	AAAA
MOTA	1364	N		A 173		24.364	25.026		_	
ATOM	1365	CA		A 173		24.253	23.680	53.620	1.00 19.91	AAAA
MOTA	1366	CB		A 173		23.342	23.699	52.397	1.00 20.86	AAAA
MOTA	1367	CG	ASP A	A 173		23.780	24.719	51.358	1.00 21.90	AAAA
ATOM	1368	OD1	ASP A	A 173		23.257	24.640	50.217	1.00 21.35	AAAA
ATOM	1369	OD2	ASP A	A 173		24.624	25.597	51.687	1.00 21.35	AAAA
ATOM	1370	C.		A 173		25.573	23.021	53.227	1.00 21.02	AAAA
	1371	ò		A 173		25.673				
ATOM			OT W	174			23.832	52.912	1.00 20.03	AAAA
ATOM	1372	N		A 174		26.579				
ATOM	1373	CA		A 174		27.870	23.277	52.553	1.00 19.72	AAAA
ATOM	1374	С	GLY A	A 174		28.537	22.680		1.00 20.27	AAAA
MOTA	1375	0	GLY A			29.110	21.599	53.711	1.00 19.77	AAAA
ATOM	1376	N	VAL F	175		28.448	23.387	54.893	1.00 21.38	AAAA
ATOM	1377	CA	VAL .			29.056	22.934	56.135	1.00 22.26	AAAA
	1378	CB	VAL ?			29.032	24.040	57.203	1.00 23.15	AAAA
MOTA			VAL A	175		29.853	23.617	58.418	1.00 22.84	AAAA
ATOM	1379	CG1	VAL	175					1.00 23.43	AAAA
ATOM	1380		VAL A	7 T/2		29.562	25.347	.56.612		
MOTA	1381	С	VAL A			28.302	21.724	56.654	1.00 23.51	AAAA
ATOM	1382	0	VAL A	175		28.893	20.803	57.210	1.00 23.74	AAAA
ATOM	1383	N	GLN A	176		26.993	21.721	56.452	1.00 24.80	AAAA
MOTA	1384	CA	GLN A			26.171	20.601	56.893	1.00 25.41	AAAA
	1385	СВ	GLN A			24.689	20.913	56.694	1.00 24.77 .	AAAA
MOTA	1386	CG	GLN A			23.799	19.735	57.036	1.00 26.23	AAAA
ATOM .	Y200	CG	GUN F				,		_ · · ·	

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Figure 19-22

							00 224	20 004	57.069	1.00 27.17	AAAA
ATOM	1387	CD	GLN	А	176		22.334	20.094			
ATOM	1388	OE1	GLN	А	176		21.902	20.879	57.911	1.00 28.24	AAAA
	1389	NE2	_				21.556	19.522	56.151	1.00 26.54	AAAA
MOTA							26.512	19.293	56.180	1.00 25.67	AAAA
MOTA	1390	С	GLN							1.00 26.98	AAAA
MOTA	1391	0	GLN	Α	176		26.789	18.285	56.820		
ATOM-	1392	N	GLU	Α	177		26.490	19.309	54.853	1.00 26.34	AAAA
	1393	CA	GLU				26.786	18.117	54.073	1.00 26.18	AAAA
MOTA							26.746	18.468	52.580	1.00 -27.13	AAAA
MOTA	1394	CB	GLU								AAAA
MOTA	1395	CG	GLU	А	177		26.769	17.269	51.628	1.00 29.77	
ATOM	1396	CD .	GLU	Α	177		26.623	17.660	50.147	1.00 31.29	AAAA
			GLU				27.655	17.935	49.500	1.00 31.35	AAAA
ATOM	1397							17.703	49.636	1.00 32.05	AAAA
MOTA	1398	OE2	GLU				25.471				AAAA
ATOM	1399	C	GLU	Α	177	•	28.160	17.556	54.460	1.00 26.38	-
ATOM	1400	0	GLU	Α	177		28.338	16.349	54.595	1.00 25.41	AAAA
		N	ALA				29.115	18.458	54.659	1.00 -27.58	AAAA
ATOM	1401						30.495	18.119	55.004	1.00 27.66	- AAAA
MOTA	1402	CA	AĻA							1.00 26.20	AAAA
MOTA	1403	CB	ALA				31.345	19.385	54.994		
ATOM	1404	Ç	ALA	Α	178		30.713	17.370	56.318	1.00 28.33	AAAA
	1405	Õ	ALA				31.685	16.626	56.439	1.00 29.44	AAAA
MOTA							29.849	17.564	57.308	1.00 28.25	AAAA
MOTA	1406	N	PHE						58.561	1.00 29.20	AAAA
MOTA	1407	CA	PHE				30.036	16.852			
MOTA	1408	CB	PHE	А	179		30.570	17.794	59.624	1.00 29.35	AAAA
	1409	CG	PHE				31.751	18.572	59.171	1.00 30.26	AAAA
MOTA			PHE				31.582	19.777	58.497	1.00 31.01	AAAA
MOTA	1410							18.069	59.339	1.00 30.37	AAAA
MOTA	1411		PHE				33.033				
MOTA	1412	CE1	PHE	А	179		32.670	20.470	57.993	1.00 31.20	AAAA
ATOM	1413	CE2	PHE	Α	179		34.133	18.749	58.840	1.00 31.74	AAAA
		CZ			179		33.950	19.960	58.161	1.00 31.81	AAAA
MOTA	1414							16.180	59.040	1.00 30.33	AAAA
ATOM	1415	С			179		28.760			1.00 31.82	AAAA
ATOM	1416	0	PHE	Α	179		28.624	15.810	60.215		
MOTA	1417	N	TYR	Α	180		27.842	15.994	58.105	1.00 29.18	AAAA
	1418	CA			180		26.564	15.379	58.389	1.00 28.99	; AAAA
ATOM					180		25.725	15.343	57.123	1.00 28.30	AAAA
MOTA	1419	CB			_				57.384	1.00 28.27	AAAA
ATOM	1420	CG			180		24.244	15.422			
MOTA	1421	CD1	TYR	Α	180		23.392	14.386	57.021	1.00 27.05	AAAA
	1422		TYR				22.029	14.491	57.197	1.00 28.51	AAAA
ATOM		CD2					23.686	16.573	57.942	1.00 29.34	AAAA
MOTA	1423							16.691	58.125	1.00 29.27	AAAA
MOTA	1424	CE2			180		22.316			1.00 29.33	AAAA
MOTA	1425	CZ	TYR	A	180		21.495	15.645	57.746		
ATOM	1426	OH	TYR	A	180		20.141	15.775	57.893	1.00 30.83	AAAA
	1427	C			180		26.673	13.970	58.940	1.00 28.79	AAAA
ATOM							25.877	13.577	59.785	1.00 28.71	AAAA
MOTA	1428	0			180			13.214	58.472	1.00 29.03	AAAA
MOTA	1429	N			181		27.662				AAAA
MOTA	1430	CA	ASP	Α	181		27.813	11.828	58.914	1.00 28.49	
	1431	CB	ASP	Α	181		28.140	10.930	57.715	1.00 27.59	AAAA
MOTA		CG			181		29.548	11 122	57.229	1.00 28.82	AAAA
MOTA	1432						29:981	12 292	57.183	1.00 29.25	AAAA
ATOM	1433		ASP				29:301	10 110		1.00 28.68	AAAA
ATOM	1434	OD2	ASP	A	181			,10 119	56.887		
ATOM	1435	С	ASP	Α	181		28.863	11.631	60.009	1.00 27.67	AAAA
	1436	0			181		29.271	10.504	60.293	1.00 27.57	AAAA
ATOM							29.308	12.713	60.628	1.00 26.27	AAAA
ATOM	1437	N			182				61.689	1.00 26.22	AAAA
ATOM	1438	CA	THR	A	182		30.284	12.544		1.00 20.22	
ATOM	1439	CB	THR	Α	182		31.670	13.118	61.317	1.00 25.92	AAAA
	1440		THR	Δ	182		32.564	12.935	62.416	1.00 25.06	AAAA
ATOM		202	min		102		31.577	14.594	60.974	1.00 25.25	AAAA
MOTA	1441		THR						62.934	1.00 25.87	AAAA
ATOM	1442	С			182		29.792	13.223			
ATOM	1443	0	THR	A	182		28.942	14.102	62.863	1.00 26.35	AAAA
	1444	Ŋ			183		30.327	12.804	64.071	1.00 25.86	AAAA
ATOM							29,953	13.370	65.355	1.00 26.12	AAAA
MOTA	1445	CA			183				66.274		AAAA
ATOM	1446	CB			183		29.468	12.260			
ATOM	1447	CG	ASP	Α	183		30.515	11.194	66.488		AAAA
	1448		ASP				31.063	10.691	65.480		AAAA
ATCM							30.781	10.852	67.657		AAAA
ATOM	1449		ASP						65.995		AAAA
ATOM	1450	C			183		31.126	14.120			
ATOM	1451	0	ASP	Α	183		31.034	14.566			AAAA
	1452	N	GT.N	A	184		32.229	14.254	65,254	1.00 26.05	AAAA
ATCM	1472										

MOTA	1453	CA	GLN	Α	184	33.381	14.983	65.766	1.00	25.85	AAAA
	2 A E A		CTN	>	104	34.674	14.510	65.095			
MOTA	1454	CB	GTM	M	184				1.00	26.30	AAAA
MOTA	1455	CG	GLN	Α	184	34.920	13.030	65.303	1.00	27.42	AAAA
ATOM	1456	CD			184	36.273	12.583	64.822	1.00	28.40	AAAA
MOTA	1457	OE1	GLN	Α	184	36.685	12.905	63.709	1.00	30.05	AAAA
	1458					36.970	11.816	65.651			
MOTA		WEZ	GLN						1.00	29.24	AAAA
MOTA	1459	С	GLN	Α	184	33.159	16.474	65.536	1.00	25.22	. AAAA
							17.316	66.220			
MOTA	1460	0			184	33.734			1.00	24.57	AAAA
MOTA	1461	N	VAL	Α	185	32.290	16.791	64.584	1.00	25.17	AAAA
						31.975	18.182	64.291			
MOTA	1462	CA			185					24.49	AAAA
ATOM	1463	CB	VAL	Α	185	32.324	18.563	62.832	1.00	23.20	AAAA
	1464	CCI	VAL	λ	105	32.045	20.060	62.599		19.72	AAAA
MOTA											
MOTA	1465	CGZ	VAL	Α	185	33. <i>777</i>	18.205	62.543	1.00	20.67	AAAA
MOTA	1466	С	VAL	Δ	185	30.494	18.421	64.501	1 00	24.81	AAAA
ATOM	1467	0	VAL	A	185	29.664	17.787	63.844	1.00	27.07	AAAA
ATOM	1468	N	PHE	Α	186	30.162	19.311	65.434	1.00	23.40	AAAA
MOTA	1469	CA			186	28.768	19.645	65.684	1.00	20.31	AAAA
ATOM	1470	CB	PHE	Α	186	28.513	19.937	67.164	1.00	19.77	AAAA
	1471	CG			186	27.057	20.037	67.500		18.55	AAAA
MOTA											
MOTA	1472	CD1	PHE	A	186	26.359	18.918	67.945	1.00	17.70	AAAA
ATOM	1473	CD2	PHE	Δ	186	26.358	21.213	67.263	1 00	17.46	AAAA
MOTA	1474	CET	PHE	A	186	24.999	18.964	68.147	1.00	17.35	AAAA
MOTA	1475	CE2	PHE	Α	186	24.997	21.271	67.459	1.00	18.83	AAAA
MOTA	1476	CZ	PHE	А	180	24.308	20.138	67.905		18.67	AAAA
MOTA	1477	С	PHE	Α	186	28.464	20.911	64.895	1.00	19.18	AAAA
	1478	0	PHE			29.079	21.940	65.129	1 00	18.82	AAAA
MOTA	_										
MOTA	1479	N	VAL	Α	187	27.520	20.834	63.964	1.00	18.34	AAAA
ATOM	1480	CA	VAL	Δ	187	27.137	21.993	63.160	3 00	16.47	AAAA
MOTA	1481	CB	VAL	А	187	27.006	21.630	61.655	1.00	14.30	AAAA
ATOM	1482	CG1	VAL	А	187	26.628	22.869	60.828	1.00	10.34	AAAA
MOTA	1483	CGZ	AYT	A	187	28.314	21.031	61.160		12.07	AAAA
MOTA	1484	С	VAL	Α	187	25.806	22.511	63.665	1.00	17.43	AAAA
	1485	0	VAL			24.852	21.746	63.792		16.95	AAAA
ATOM											
ATOM	1486	N	LEU	Α	188	25.763	23.809	63.960	1.00	18.66	AAAA
ATOM	1487	CA	LEU	Α	188	24.555	24.507	64.460	1.00	20.51	AAAA
MOTA	1488	CB	LEU	A	188	24.752	24.995	65.914	1.00	21.24	AAAA
ATOM	1489	CG	LEU	А	188	23.702	26.019	66.395	1.00	20.80	AAAA
			LEU			22.365	25.323	66.493		19.77	
MOTA	1490										AAAA
ATOM	1491	CD2	LEU	A	188	24.085	26.627	67.750	1.00	20.63	AAAA
ATOM	1492	C	LEU	Δ	199	24.297	25.735	63.591	1 00	20.41	AAAA
ATOM	1493	0	LEU	А	188	25.223	26.484	63.288	1.00	21.86	AAAA
ATOM	1494	N	SER	Α	189	23.049	25.987	63.233	1.00	19.32	AAAA
ATOM	1495	CA	SER	A	193	22.786	27.130	62.381	1.00	18.06	AAAA
MOTA	1496	CB	SER	А	189	22.970	26.715	60.906	1.00	18.54	AAAA
		OG				22.559	27.731	59.998		17.47	AAAA
MOTA	1497		SER								
MOT A	1498	С	SER	А	189	21.418	27.751	62.554	1.00	17.90	AAAA
MOLA	1499	0	SER	Δ	189	20.404	27.051	62.540	1 00	19.54	AAAA
A_OM	1500	N	LEU	A	190	21.386	29.067	62.722	1.00	16.97	AAAA
ATOM	1501	CA	LEU	Α	190	20.117	29.772	62.797	1.00	18.49	AAAA
ATOM	1502	CB	LEU	A	190	20.097	30.865	63.886		17.78	AAAA
ATOM	1503	CG	LEU	Α	190	20.534	30.600	65.337	1.00	17.10	AAAA
	-		LEU			19.643	31.406			15.50	
ATOM	1504										AAAA
MOTA	1505	CD2	LEU	A	190	20.455	29.147	65.686	1.00	15.15	AAAA
	1506	C	LEU	Δ	190	20.111	30.408	61.416 .	1 00	19.35	AAAA
ATOM											
ATOM	1507	0	LEU			21.136	30.891	60.967		19.75	AAAA
ATOM	1508	N	HIS	Α	191	18.975	30.397	60.736	1.00	21.75	AAAA
								59.383			
MOTA	1509	CA	HIS			18.897	30.955			23.55	AAAA
ATOM	1510	CB	HIS .	Α	191	19.626	30.013	58.426	1.00	23.63	AAAA
							28.597	58.533		24.26	
MOTA	1511	CG	HIS			19.157					AAAA
MOTA	1512	CD2	HIS	Α	191	19.770	27.485	59.009	1.00	23.78	AAAA
			HIS .			17.869	28.217	58.217		24.73	AAAA
MOTA	1513										
MOTA	1514	CE1	HIS.	Α	191	17.709	26.935	58.491		23.90	AAAA
	1515		HIS.			18.849	26.467	58.973	1.00	24.51	AAAA
ATOM											
MOTA	1516	С	HIS .			17.446	31.119	58.926		24.10	AAAA
ATOM	1517	0	HIS.	Α	191	16.519	30.658	59.596	1,00	24.94	AAAA
							31.789	57.794		24.33	AAAA
MOTA	1518	N	GLN .	~	134	17.249	31./03	J / J 4	1.00	24.73	

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Figure 19-24

MOTA	1519	CA	GLN A	192	15.899	31.959	57.269	1.00 25.77	AAAA
								1.00 26.51	AAAA
ATOM	1520	CB	GŢN A		15.881	32.896	56.060		
ATOM	1521	CG	GLN A	192	16.467	34.271	56.325	1.00 26.99	AAAA
ATOM	1522	CD	GLN A	192	16.581	35.076	55.062	1.00 27.98	AAAA
ATOM	1523	OE1	GLN A	192	15,583	35.496	54.493	1.00 30.48	AAAA
			GLN A		17.802	35.274	54.595	1.00 29.04	AAAA
MOTA	1524		_						
ATOM	1525	С	GLN A		15.463	30.573	56.832	1.00 25.77	AAAA
MOTA	1526	0	GLN A	192	16.211	29.865	56.169	1.00 26.73	AAAA
ATOM	1527	N	SER A	193	14.259	30.184	57.214	1.00 25.48	AAAA
	1528	CA	SER A		13.750	28.877	56.863	1.00 24.51	AAAA
MOTA								1.00 23.77	AAAA
ATOM	1529	CB	SER A		12.288	28.788	57.286		
ATOM	1530	OG	SER A	193	11.753	27.517	57.010	1.00 24.81	AAAA
MOTA	1531	С	SER A	193	13.906	28.597	55.361	1.00 24.53	AAAA
MOTA	1532	0	SER A	193	13.736	29.479	54.522	1.00 22.32	AAAA
		N	PRO A		14.226	27.348	55.007	1.00 25.69	AAAA
MOTA	1533						55.862	1.00 25.02	
MOTA	1534	CD	PRO A		14.411	26.167			AAAA
ATOM	1535	CA	PRO A	194	14.399	26.976	53.604	1.00 27.05	AAAA
MOTA	1536	CB	PRO A	194	14.906	25.535	53.697	1.00 26.30	AAAA
ATOM	1537	CG	PRO A	194	15.479	25.466	55.124	1.00 26.44	AAAA
			PRO A		13.076	27.057	52.849	1.00 27.79	AAAA
MOTA	1538	C							
MOTA	1539	0	PRO A	194	13.066	27.057	51.625	1.00 28.82	AAAA
ATOM	1540	N	GLU A	195	11.966	27.133	53.582	1.00 28.29	AAAA
MOTA	1541	CA	GLU A	195	10.656	27.187	52.950	1.00 29.08	AAAA
	1542	CB	GLU A		9.534	27.030	54.001	1.00 31.08	AAAA
MOTA					9.070	28.294	54.722	1.00 35.07	AAAA
MOTA	1543	CG	GLU A						
MOTA	1544		GLU A		7.850	28.980	54.064	1.00 38.05	AAAA
ATOM	1545	OE1	GLU A	195	7.389	30.017	54.601	1.00 38.80	AAAA
ATOM	1546		GLU A		7.342	28.487	53.024	1.00 39.20	AAAA
	1547	C	GLU A		10.483	28.471	52.150	1.00 28.05	AAAA
MOTA					9.722	28.512	51.189	1.00 28.57	AAAA
MOTA	1548	0	GLU A						
MOTA	1549	N	TYR A	196	11.223	29.510	52.514	1.00 27.39	AAAA
ATOM	1550	CA	TYR A	196	11.108	30.769	51.802	1.00 25.80	AAAA
ATOM	1551	CB	TYR A	196	10.275	31.743	52.645	1.00 24.97	AAAA
	1552	CG	TYR A		10.971	32,281	53.868	1.00 23.41	AAAA
ATOM					11.911	33.306	53.765	1.00 23.99	AAAA
ATOM	1553		TYR A						AAAA
ATOM	1554	CEl			12.559	33.805	54.892	1.00 23.44	
ATOM	1555	CD2	TYR A	196	10.697	31.768	55.126	1.00 23.24	AAAA
MOTA	1556	CE2	TYR A	196	11.336	32.256	56.254	1.00 23.93	AAAA
ATOM	1557	CZ	TYR A	196	12.265	33.270	56.133	1.00 24.07	AAAA
		OH	TYR A		12.913	33.731	57.247	1.00 25.06	AAAA
ATOM	1558						51.411	1.00 24.97	AAAA
MOTA	1559	С	TYR A		12.450	31.406			
ATOM	1560	0	TYR A		12.475	32.495	50.840	1.00 25.14	AAAA
MOTA	1561	N	ALA A	197	13.563	30.737	51.686	1.00 23.81	AAAA
ATOM	1562	CA	ALA A	197	14.855	31.330	51.337	1.00 23.32	AAAA
		CB	ALA A		15.350	32.220	52.488	1.00 23.33	AAAA
ATOM	1563					30.356	50.957	1.00 22.74	AAAA
- MOTA	1564	С	ALA A		15.952	_			
MOTA	1565		ALA A		15.951		51.37	1.00 22.47	AAAA
ATOM -	1566	N	.PHE A	- 198	16, 900	30.852	50.16	1.00 23.23	AAAA
ATOM	1567	CA	PHE A	198	18.062	30.081	49.741	1.00 23.68	AAAA
	1568	CB	PHE A		19.083	31.006	49.069	1.00 23.33	AAAA
MOTA							48.464	1.00 22.98	AAAA
ATOM	1569	CG	PHE A		20.250	30.280			
MOTA	1570	CD1	PHE A	198	20.151	29.713	47.203	1.00 22.75	AAAA
ATOM	1571	CD2	PHE A	198	21.436	30.127	49.175	1.00 23.32	AAAA
MOTA	1572	CEI	PHE A	198	21.207	29.003	46.645	1.00 22.13	AAAA
		CES	PHE A	109	22.512	29.408	48.622	1.00 22.83	AAAA
ATOM	1573		rne A	100		28.849	47.351	1.00 22.55	· AAAA
ATOM	1574	CZ	PHE A	T30	22.386				
MOTA	1575	С	PHE A		18.689	29.490	51.008	1.00 23.69	AAAA
ATOM	1576	0	PHE A	198	18.802	30.171	52.012	1.00 22.85	AAAA
	1577	N	PRO A	199	19.166	28.236	50.954	1.00 23.96	AAAA
MOTA			PRO A	1.0.0	19.833	27.639	52.123	1.00 24.26	AAAA
MOTA	1578	CD						1.00 24.70	AAAA
ATOM	1579	CA	PRO A	199	19.199	27.286	49.837		
ATOM	1580	CB	PRO A	199	20.163	26.222	50.357		AAAA
MOTA	1581	CG	PRO A	199	19.797	26.162	51.782	1.00 23.21	AAAA
	1582	c	PRO A		17.885	26.679	49.326	1.00 25.22	AAAA
MOTA			PRO A	100	17.866	26.145	48.215	1.00 26.24	AAAA
ATOM	1583	0	PRU A	200			50.116	1.00 25.09	AAAA
ATOM	1584	N	PHE A	200	16.811	26.756	20.110	1.00 23.09	MANA

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Figure 19-25

2001	1 5 0 5	CA	PHE A	200		15.497	26.190	49.763	1.00 26.2 9	AAAA
MOTA	1585		FIII A	200		15.064	26.567	48.340	1.00 25.65	AAAA
ATOM	1586	CB	PHE A							AAAA
MOTA	1587	CG	PHE A	200		14.863	28.035	48.122	1.00 24.65	
	1588	CD1	PHE A	200	•	15.806	28.781	47.439	1.00 24.42	AAAA
ATOM		CDI	PHE A	200		13.735	28.671	48.608	1.00 23.79	AAAA
MOTA	1589	CDZ	PHE A	200				47.246	1.00 24.41	AAAA
ATOM_	1590	CE1	PHE A	200.		15.631	30.125			
MOTA	1591	CE2	PHE A	200		13.552	30.035	48.418	1.00 24.94	, AAAA
	1592	cz	PHE A			14.499	30.760	47.738	1.00 24.57	AAAA.
MOTA			_			15.415	24.656	49.863	1.00 28.54	AAAA
MOTA	1593	С	PHE A							AAAA
MOTA	1594	0	PHE A	200		14.386	24.096	50.251	1.00 28.76	
	1595	N	GLU A			16.499	23.981	49.504	1.00 29.67	AAAA
MOTA						16.539	22.528	49.524	1.00 31.88	AAAA
MOTA	1596	CA	GLU A					48.392	1.00 32.71	AAAA
ATOM	1597	CB	GLU A	201	•	17.434	22.045			
ATOM	1598	CG	GLU A	201		16.897	22.415	47.017	1.00 34.87	AAAA
		CD	GLU A			17.898	22.147	45.912	1.00 35.14	AAAA
MOTA	1599					18.299	20.982	45.735	1.00 36.09	- AAAA
MOTĄ	1.600		GLU _. A			10.233		45.221	1.00 36.30	AAAA
ATOM	1601	OE2	GLU A	201		18.286				
MOTA	1602	С	GLU A	201		16.997	21.894	50.835	1.00 32.77	AAAA
			GLU A			16.806	20.690	51.046	1.00 33.44	AAAA
MOTA	1603	0					22.690	51.711	1.00 32.31	AAAA
MOTA	1604	N	LYS A			17.599				AAAA
ATOM	1605	CA	LYS A	202		18.101	22.168	52.974	1.00 32.09	
	1606	СВ	LYS A			19.565	21.750	52.811	1.00 33.02	AAAA
MOTA						19.836	20.847	51.623	1.00 34.95	AAAA
MOTA	1607	CG	LYS A					51.436	1.00 37.92	AAAA
MOTA	1608	CD	LYS A			21.334	20.619			
MOTA	1609	CE	LYS A	202		21.655	19.804	50.169	1.00 39.19	AAAA
		NZ	LYS A			23.120	19.522	49.988	1.00 38.58	AAAA
MOTA	1610					17.995	23.241	54.037	1.00 30.85	AAAA
MOTA	1611	С	LYS A						1.00 30.49	AAAA
MOTA	1612	0	LYS A	202		17.706	24.389	53.739		
MOTA	1613	N	GLY A	203		18.238	22.867		1.00 30.81	AAAA
			GLY A			18.159	23.831	56.356	1.00 30.86	AAAA
MOTA	1614	CA					23.578	57.280	1.00 30.84	AAAA
MOTA	1615	С	GLY A			16.991				AAAA
MOTA	1616	0	GLY A	203		16.828	24.285	58.272	1.00 31.58	
	1617	N	PHE A	204		16.182	22.570	56.965	1.00 30.54	AAAA
MOTA			PHE A			15.025	22.241	57.797	1.00 30.51	AAAA
MOTA	1618	CA					21.317		1.00 29.06	AAAA
MOTA	1619	CB	PHE A			14.061				AAAA
ATOM	1620	CG	PHE A	204		13.524	21.890	55.787		
	1621	ന1	PHE A	204		14.222	21.762	54.601	1.00 26.52	AAAA
MOTA			PHE A			12.307	22.548	55.779	1.00 26.50	AAAA
MOTA	1622						22.276	53.420	1.00 26.44	AAAA
ATOM	1623		PHE A			13.713				AAAA
ATOM	1624	CE2	PHE A	204		11.786	23.069	54.600	1.00 26.69	
•	1625	CZ	PHE A	204		12.490	22.931	53.416	1.00 25.65	AAAA
MOTA						15.401	21.590	59.127	1.00 30.87	AAAA
MOTA	1626	С	PHE A				20.875	59.228	1.00 31.12	AAAA
MOTA	1627	0	PHE A			16.395				AAAA
ATOM	1628	N	LEU A	205		14.580	21.844		1.00 31.22	
	1629	CA	LEU A	205		14.782	21.329	61.489	1.00 31.43	AAAA
ATOM			LEU A			13.575	21.691	62.357	1.00 31.42	AAAA
ATOM	1630					13.575	21.078		1.00 31.76	AAAA
MOTA	1631	CG	LEU A			13.603				AAAA
ATOM	1632	CD1	LEU A	235		14.894	21.492		1.00 32.36	
	1633		LEU A			12.379	21.516	64.536	1.00 31.31	AAAA
MOTA						15.026	19.829		1.00 31.35	AAAA
MOTA	1634	С	LEU A						1.00 31.40	AAAA
MOTA	1635	0	LEU A	205		15.714	19.392			
ATOM	1636	N	GLU A	206		14.448	19.059	60.707	1.00 31.79	AAAA
		CA	GLU A			14.509	17.603	60.706	1.00 32.08	AAAA
ATOM	1637					13,485	17.054	59.716	1.00 33.18	AAAA
MOTA	1638	CB	GLU A							AAAA
ATOM	1639	CG	GLU A			12.069	17.651		1.00 34.20	
	1640	CD	GLU A			11.973	19.136	59.453	1.00 33.44	AAAA
MOTA						10.854	19.675		1.00 33.32	AAAA
MOTA	1641		GLU A						1.00 35.12	AAAA
MOTA	1642	OE2	GLU A			13.005	19.777			
	1643	С	GLU A			15.882	17.045	60.363	1.00 32.34	AAAA
ATOM			GLU A			16.209	15.909		1.00 31.83	AAAA
MOTA	1644	0					17.847			AAAA
MOTA	1645	N	GLU ?			16.680				AAAA
MOTA	1646	CA	GLU A	207		18.017	17.431		1.00 31.67	
		CB	GLU A			18.552	18.385	58.238	1.00 30.39	AAAA
MOTA	1647					17.768	18.316			AAAA
MOTA	1648	CG	GLU A							AAAA
MOTA	1649	CD	GLU A	207		17.953	19.547			
MOTA	1650		GLU A			19.108	19.991	55.971	1.00 30.31	AAAA
7.17 114		~~~	:	. -		_ , .		_		

3 mov.	1651	OES	GLU A	207	16.947	20.070	55.604	1.00 30.76	AAAA
ATOM					18.879	17.433	60.537	1.00 32.04	AAAA
ATOM	1652		GLU A				60.910	1.00 31.57	AAAA
MOTA	1653	0	GLU A	207	19.472	18.448			
MOTA	1654	N	ILE A	208	18.935	16.272	61.178	1.00 32.57	AAAA
ATOM	1655	CA	ILE A		19.674	16.111	62.408	1.00 33.37	AAAA
		СВ	ILE A		18.709	15.647	63.519	1.00 33.65	AAAA
MOTA	1656					15.380	64.806	1.00 34.11	AAAA
MOTA	1657		ILE A		19.443		_	1.00 33.94	AAAA
MOTA	1658	CG1	ILE A	208	17.673	16.742	63.757		
ATOM	1659	CD1	ILE A.	208	16.628	16.386	64.794	1.00 37.00	AAAA
MOTA	1660	C	ILE A		20.863	15.174	62.280	1.00 34.00	AAAA
			ILE A		21.506	14.829	63.265	1.00 34.40	AAAA
ATOM	1661	0				14.768	61.062	1.00 34.64	AAAA
MOTA	1662	N	GLY A		21.177			1.00 35.55	AAAA ·
MOTA	1663	CA	GLY A	209	22.321	13.903	60.913		
ATOM	1664	С	GLY A	209	22.164	12.671	60.057	1.00 36.80	AAAA
	1665	ō	GLY A		21.148	12.461	59.400	1.00 37.32	AAAA
MOTA			GLU A		23.199	11.836	60.100	1.00 37.78	AAAA
MOTA	1666	N				10.621	59.315	1.00 38.04	AAAA
MOTA	1667	CA	GLU A		23.256			1.00 38.54	AAAA
MOTA	1668	CB	GLU A	210	23.600	11.013	57.892		
MOTA	1669	CG	GLU A	210	23.469	9.960	56.858	1.00 38.99	AAAA
	1670	CD	GLU A		24.118	10.412	55.580	1.00 40.10	AAAA
ATOM			GLU A		25.365	10.437	55.555	1.00 40.86	AAAA
MOTA	1671					10.767	54.619	1.00 40.41	AAAA
MOTA	1672	OE2			23.396			1.00 37.98	AAAA
ATOM	1673	С	GLU A	210	24.377	9.770	59.894		
ATOM	1674	0	GLU A	210	25.498	10.244	60.041	1.00 38.52	AAAA
	1675	N	GLY A	211	24.085	8.517	60.220	1.00 38.02	AAAA
MOTA			GLY A		25.116	7.654	60.770	1.00 38.09	AAAA
atom	1676	CA			25.542	8.075	62.164	1.00 38.26	AAAA
ATOM	1677	С	GLY A				62.977	1.00 37.82	AAAA
MOTA	1678	0	GLY A	211	24.697	8.443			AAAA
ATOM	1679	N	LYS A	212	26.848	8.030	62.434	1.00 38.20	
ATOM	1680	CA	LYS A	212	27.396	8.399	63.743	1.00 37.56	AAAA
		CB	LYS A		28.921	8.209	63.766	1.00 38.86	AAAA
MOTA	1681				29.416	6.810	63.385	1.00 40.93	AAAA
MOTA	1682	CG	LYS A				64.405	1.00 42.04	AAAA
ATOM	1683	CD	LYS A		29.001	5.746		1.00 42.80	AAAA
MOTA	1684	CE	LYS A	212	29.251	4.318	63.891		
ATOM	1685	NZ	LYS A	212	30.673	4.002	63.562	1.00 42.32	AAAA
	1686	C	LYS A		27.093	9.859	64.054	1.00 37.08	AAAA
MOTA			LYS A		27.075	10.269	65.218	1.00 36.94	AAAA
ATOM	1687	0				10.636	63.002	1.00 35.41	AAAA
MOTA	1688	N	GLY A		26.854		63.170	1.00 34.24	AAAA
ATOM	1689	CA	GLY A		26.592	12.054			AAAA
ATOM	1690	С	GLY A	213	25.163	12.438	63.470	1.00 33.27	
ATOM	1691	0	GLY A	213	24.861	13.611	63.666	1.00 33.29	AAAA
		N	LYS A		24.280	11.451	63.512	1.00 31.79	AAAA
MOTA	1692				22.883	11.710	63.794	1.00 30.47	AAAA
MOTA	1693	CA	LYS A			10.396	63.737	1.00 30.35	AAAA
ATOM	1694	CB	LYS Y		22.111		63.280	1.00 30.45	AAAA
MOTA	1695	CG	LYS A	214	20.676	10.552			
ATOM	1696	CD	LYS A	214	20.141	9.241	62.759	1.00 29.75	AAAA
	1697	CE	LYS A	214	18.737	9.400	62.229	1.00 30.19	AAAA.
MOTA		NZ	LYS A		18.179	8.138	61.671	1.00 31.35	AAAA
ATOM.	1698					12.374	65.168	1.00 30.31	AAAA
MOTA	1699	C	LYS A		22.778	11.814	66.177	1.00 30.44	AAAA
ATOM	1700	0	LYS A		23.193			1.00 29.95	AAAA
ATOM	1701	N	GLY A	215	22.243	13.590	65.192		
ATOM	1702	CA	GLY A	215	22.128	14.325	66.437	1.00 29.16	AAAA
	1703	c	GLY A		23.222	15.379	66.582	1.00 28.54	AAAA
MOTA					23.306	16.061	67.602	1.00 28.27	AAAA
MOTA	1704	0	GLY A			15.521	65.561	1.00 27.35	AAAA
ATOM	1705	Ņ	TYR A	. 216	24.063			1.00 27.08	AAAA
ATOM	1706	CA	TYR A	. 216	25.150	16.497	65.616		
MOTA	1707	·CB	TYR A		26.516	15.800	65.531	1.00 28.38	AAAA
	1708	CG	TYR A		26.786	14.966	66.757	1.00 30.21	AAAA
ATOM			TYR A		26.138	13.735	66.955	1.00 29.81	AAAA
atom	1709				26.311	13.014	68.138	1.00 30.03	AAAA
ATOM	1710		TYR A					1.00 29.71	AAAA
ATOM	1711	CD2	TYR A	216	27.619				
ATOM	1712	CE2		216	27.798	14.741		1.00 29.96	AAAA
		CZ	TYR A		27.143	13.528	69.138	1.00 30.84	AAAA
MOTA	1713		TYR A		27.297			1.00 31.14	AAAA
atcm	1714	OH							
MOTA	1715	C	TYR A		25.055			1.00 26.38	
ATCM	1716	0	TYR A	216	26.046	18.240	204.243	1.00 20.30	
			•						-

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Figure 19-27

ATOM	1717	N	ASN A	217	23.845	17.791	64.076	1.00 23.55	AAAA
MOTA	1718	CA	ASN A		23.549	18.830	63.119	1.00 21.52	AAAA
MOTA	1719	CB	ASN A		23.431	18.282	61.699	1.00 20.64	AAAA
MOTA	1720	CG	ASN A		23.202	19.386	60.669 60.499	1.00 20.29 1.00 18.03	AAAA AAAA
MOTA	1721		ASN A		22.089 24.274	19.888 19.790	60.499	1.00 19.95	AAAA
MOTA	1722		ASN A		24.274	19.790	63.605	1.00 21.64	AAAA
ATOM	1723 1724	.O	ASN A		21.263	18.576	63.757	1.00 20.34	AAAA
MOTA MOTA	1725	N	LEU A		22.165	20.647	63.873	1.00 22.22	AAAA
ATOM	1726	CA	LEU A		20.960	21.282	64.388	1.00 22.03	AAAA
ATOM	1727	CB	LEU A	218	21.195	21.711	65.840	1.00 20.97	AAAA
ATOM	1728	CG	LEU A		20.051	21.838	66.841	1.00 20.94	AAAA
MOTA	1729		LEU A		20.513	22.744	67.936	1.00 20.31 1.00 21.27	AAAA AAAA
MOTA	1730		LEU A		18.818 20.669	22.412 22.513	66.227 63.547	1.00 21.27	AAAA
ATOM	1731	C	LEU A		21.451	23.454	63.557	1.00 22.70	AAAA
MOTA	1732 1733	о И	ASN A		19.564	22.491	62.808	1.00 24.00	AAAA
ATOM ATOM	1734	CA	ASN A		19.166	23.626	61.990	1.00 25.33	AAAA
MOTA	1735	CB	ASN A		18.656	23.190	60.614	1.00 26.94	AAAA
MOTA	1736	CG	ASN. A	219	19.737	22.601	59.749	1.00 26.68	AAAA
MOTA	1737		ASN A		20.812	23.169	59.626	1.00 28.06	AAAA
MOTA	1738		ASN A		19.446	21.471	59.117	1.00 26.26	AAAA
MOTA	1739	C	ASN A		18.046	24.345	62.710 63.210	1.00 25.69 1.00 27.51	AAAA AAAA
MOTA	1740	0	ASN A ILE A		17.118 18.122	23.706 25.667	62.753	1.00 27.50	AAAA
MOTA	1741 1742	N CA	ILE A		17.107	26.457	63.428	1.00 25.87	AAAA
MOTA MOTA	1743	CB	ILE A		17.733	27.331	64.557	1.00 25.04	AAAA
MOTA	1744		ILE A		16.654	28.152	65.227	1.00 25.24	AAAA
MOTA	1745	CG1	ILE A	220	18.460	26.447	65.584	1.00 24.07	AAAA
MOTA	1746	CD1	ILE A		17.557	25.502	66.378	1.00 22.28	AAAA
ATOM	1747	С	ILE A		16.430	27.370	62.414	1.00 26.20 1.00 25.35	AAAA AAAA
MOTA	1748	0	ILE A		16.801 15.421	28.534 26.850	62.265 61.704	1.00 25.33	AAAA
MOTA	1749	N CD	PRO A		14.840	25.501	61.778	1.00 27.17	AAAA
MOTA MOTA	1750 1751	CA	PRO A		14.706	27.640	60.703	1.00 27.67	AAAA
MOTA	1752	CB	PRO A		13.771	26.613	60.064	1.00 26.81	AAAA
ATOM	1753	CG	PRO A	221	14.473	25.293	60.346	1.00 27.36	AAAA
ATOM	1754	C	PRO A		13.944	28.763	61.390	1.00 28.61	AAAA
ATOM	1755	0	PRO A		13.218	28.515	62.363 60.900	1.00 29.91 1.00 28.15	AAAA AAAA
ATOM	1756	N	LEU A		14.100 13.408	29.990 31.117	,	1.00 28.48	AAAA
MOTA	1757	CA	LEU A		14.431	32.041	62.191	1.00 28.69	AAAA
MOTA	1758 1759	CB CG	LEU A		15.187	31.394	63.371	1.00 28.67	AAAA
MOTA MOTA	1760		LEU A		16.304	32.300	63.837	1.00 28.62	AAAA
MOTA	1761		LEU A		14.231	31.106	64.527	1.00 27.65	'AAAA
ATOM	1762	C	LEU A		. 12 26	31.882	60.518	1.00 28.44	AAAA
MOTA	1763	0	LEU A		12.;18	31.958	59.325	1.00 27.90 1.00 28.79	AAAA AAAA
MOTA	1764	N	PRO A		1113	32.441 32.357	61.009 62.410	1.00 29.20	AAAA
ATOM	1765	CD	PRO A		10.966 10.437	33.202	60.227	1.00 29.36	AAAA
MOTA	1766 1767	CA CB	PRO A		9.256	33.287	61.183	1.00 28.98	AAAA
ATOM	1768	CG	PRO A	223	9.965	33.502	62.485	1.00 28.68	AAAA
MOTA MOTA	1769	c	PRO A	223	10.890	34.585	59.753	1.00 30.15	AAAA
ATOM	1770	Ō	PRO A		11.864	35.152	60.253	1.00 30.18	AAAA
MOTA	1771	N	LYS A	224	10.150	35.112	58.781	1.00 30.50	AAAA
MOTA	1772	CA	LYS A		10.398	36.422	58.213	1.00 29.92	
MOTA	1773	CB	LYS A		9.491	36.661	57.008 55.893	1.00 30.57	AAAA AAAA
MOTA	1774	CG	LYS A		9.588	35.676 36.087	54.798	1.00 30.00	AAAA
MOTA	1775	CD	LYS A		8.640 8.575	35.051	53.705	1.00 32.15	AAAA
MOTA	1776	CE NZ	LYS A		7.628	35.476	52.648	1.00 32.75	AAAA
ATOM	1777 1778	C	LYS A	224	10.050	37.468	59.260	1.00 29.75	AAAA
MOTA MOTA	1779	Ö	LYS A		9.308	37.193	60.196	1.00 29.84	AAAA
ATOM	1780	N	GLY A	225	10.555	38.678	59.079	1.00 29.39	AAAA
MOTA	1781	CA	GLY A	225	10.261	39.730	60.031	1.00 29.87	AAAA
MOTA	1782	С	GLY A	225	10.809	39.447	61.415	1.00 29.85	AAAA

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Figure 19-28

		_		205		10 271	40 0E1	62.392	1.00 29.85	AAAA
ATOM	1783	0	GLY A			10.371	40.051			
MOTA	1784	N	LEU A	226		11.775	3.8.536	61.499	1.00 29.50	AAAA
ATOM	1785	CA	LEU A	226		12.374	38.175	62.778	1.00 29.80	AAAA
		CB	LEU A			13.513	37.170	62.570	1.00 28.81	AAAA
ATOM	1786	-						63.820	1.00 27.29	AAAA
ATOM	1787	CG	LEU A			14.097	36.514			
ATOM	1788	CD1	LEU A	226		13.132	35.452	64.275	1.00 26.06	AAAA
	1789	CD2	LEU A	226		15.455	35.888	63.538	1.00 27.03	AAAA
MOTA			LEU A			12.936	39.428	63.448	1.00 30.68	AAAA
ATOM	1790	C							1.00 30.57	AAAA
MOTA	1791	0	LEU A	226		13.636	40:217	62.804		
ATOM	1792	N	ASN A	227		12.624	39.617	64.729	1.00 31.46	AAAA
	1793	CA		227		13.139	40.769	65.469	1.00 32.06	AAAA
ATOM				227		12.012	41.507	66.217	1.00 31.74	AAAA
MOTA	1794	CB		-						AAAA
ATOM	1795	CG		3 227		11.291	40.630	67.234	1.00 32.07	
MOTA	1796	OD1	ASN A	A 227		11.914	40.017	68.104	1.00 31.61	AAAA
	1797		ASN A			9.962	40.592	67.141	1.00 31.59	AAAA
MOTA							40.334	66.444-	1.00 32.45	AAAA
MOTA	1798	С		3 227		14.225				AAAA
ATOM	1799	0	ASN A	A 227		14.413	39.140	66.688	1.00 32.78	
MOTA	1800	N	ASP A	A 228		14.943	41.297	67.002	1.00 33.32	AAAA
		CA		A 228		16,017	40.976	67.928	1.00 34.75	AAAA
MOTA	1801								1.00 36.77	AAAA
MOTA	1802	CB		A 228		16.508	42.233			
MOTA	1803	CG	ASP A	A 228		17.154	43.238	67.714	1.00 37.28	AAAA
	1804		ASP A	A 228		17.662	42.816	66.652	1.00 37.78	AAAA
ATOM			ASP			17.180	44.443	68.054	1.00 37.21	AAAA
MOTA	1805							68.964	1.00 34.93	AAAA
MOTA	1806	С		A 228		15.707	39.892			
ATOM	1807	Ο.	ASP A	A 228		16.448	38.919	69.056	1.00 36.92	AAAA
	1808	N .	ASN A	A 229		14.635	40.054	69.741	1.00 33.90	AAAA
MOTA				A 229		14.268	39.079	70.775	1.00 33.01	AAAA
MOTA	1809	CA						71.455	1.00 33.79	AAAA
ATOM	1810	CB		A 229		12.965	39.481			
MOTA	1811	CG	ASN :	A 229		13.131	40.663	72.369	1.00 34.04	AAAA
ATOM -			ASN .	A 229		13.783	40.564	73.405	1.00 34.25	AAAA
			ASN .			12.550	41.797	71.988	1.00 34.05	AAAA
MOTA	1813							70.276	1.00 32.98	AAAA
ATOM	1814	C		A 229		14.114	37.656			
MOTA	1815	0	ASN .	A 229		14.529	36.697		1.00 32.77	AAAA
	1816	N	GLII	A 230		13.496	37.523	69.108	1.00 32.02	AAAA
ATOM				A 230		13.277	36.227	68.516	1.00 30.72	AAAA
MOTA	1817	CA						67.272	1.00 31.38	AAAA
ATOM	1818	CB		A 230		12.399	36.375			AAAA
MOTA	1819	CG	GLU .	A 230		11.006	36.896	67.583	1.00 31.02	
ATOM	1820	CD	GLU .	A 230		10.175	37.187	66.350	1.00 31.52	AAAA
				A 230		10.644	37.970	65.497	1.00 31.89	AAAA
MOTA	1821						36.655	66.241	1.00 31.04	AAAA
MOTA	1822	OE2		A 230		9.047				
ATOM	1823	С	GLU .	A 230		14.628	35.622	68.180	1.00 30.79	AAAA
MOTA	1824	0	GLU	A 230		14.905	34.465	68.512	1.00 31.05	AAAA
		N		A 231	•	15.490	36.412	67.553	1.00 30.05	AAAA
MOTA	1825						35.920	67.191	1.00 28.94	AAAA
MOTA	1826	CA		A 231		16.811				AAAA
ATOM	1827	CB	PHE	A 231		17.632	37.015	66.528	1.00 29.33	
ATOM	1828	CG	PHE	A 231		18.949	36.537	65.972	1.00 28.79	AAAA
	1829		DUE	A 231		18.982	35.585	64.957	1.00 28.93	AAAA
Atom						20.152	37.067	66.436	1.00 28.55	AAAA
MOTA	1830			A 231					1.00 28.32	AAAA
MOTA	1831	CE1	PHE	A 231		20.195	35.160	64.397		
ATOM	1832	CE2	PHE	A 231		21.376	36.657	65.888	1.00 28.97	AAAA
				A 231		21.397	35.695	64.860	1.00 28.81	AAAA
ATOM	1833		FUE	n 231		17.559	35.443	68.413	1.00 28.25	AAAA
MOTA	1834	С		A 231					1 00 23 03	
ATOM	1835	0	PHE	A 231		17.999	34.302	68.485	1.00 27.97	AAAA
	1836	N		A 232		17.691	36.329	-69.384	1.00 27.93	AAAA
MOTA				A 232		18.425	36.003	70.590	1.00 27.93	AAAA
MOTA	1837	CA								AAAA
MOTE	1838	CB		A 232		18.521	37.234	71.484		
ATOM	1839	CG	LEU	A 232		19.220	38.379	70.747		AAAA
	1840			A 232		19.203	39.629	71.587	1.00 27.57	AAAA
ATOM		227	1 577	2 222		20.639	37.955	70.387	1.00 27.76	AAAA
ATOM	1841			A 232			_			AAAA
ATOM	1842	С	LEU	A 232		17.815	34.851			
MOTA	1843	0	LEU	A 232		18.526	34.061			AAAA
	1844	N		A 233		16.495	34.758	71.298	1.00 28.81	AAAA
ATCM			7 1177	× 222		15.786	33.685			AAAA
MOTE	1845	CA	PHE	A 233				•		AAAA
ATOM	1846	CB		A 233		14.278	33.837			
	1847	CG		A 233		13.465	32.710	72.308	1.00 32.38	AAAA
ATOM				A 233		13.257	32.599		1.00 33.66	AAAA
MOTA	1848	CDT	LUC	n LJJ			,,,,,,			

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Figure 19-29

MOTA	1849	CD2	PHE A	233		12.928	31.741	71.467	1.00 33.51	AAAA
ATOM	1850	CE1	PHE A	233		12.518	31.537	74.201	1.00 35.10	AAAA
ATOM	1851		PHE A			12.193	30.677	71.975	1.00 34.21	AAAA
							30.572	73.344	1.00 35.23	AAAA
MOTA	1852	CZ	PHE A			11.986				
MOTA	1853	С	PHE A	233		16.219	32.301	71.483	1.00 30.55	AAAA
ATOM -	1854	0	PHE A	233		16.438	31.391	72.280	1.00 30.65	AAAA
ATOM	1855	N	ALA A	234		16.317	32.151	70.165	1.00 30.21	AAAA
			ALA A			16.698	30.892	69.549	1.00 28.97	AAAA
MOTA	1856	CA								
MOTA	1857	CB	ALA A	234		16.398	30.942		1.00 30.40	AAAA
MOTA	1858	С	ALA A	234		18.169	30.571	69.761	1.00 28.27	AAAA
ATOM	1859	0	ALA A	234		18.564	29.401	69.830	1.00 26.6 9	AAAA
						18.978	31.614	69.855	1.00 27.56	AAAA
МОТА	1860	N	LEU A	-						
MOTA	1861	CA	LEU A	235	•	20.402	31.427		1.00 29.17	AAAA
MOTA	1862	CB	LEU A	235		21.126	32.767	69.989	1.00 29.04	AAAA
MOTA	1863	CG	LEU A	235		22.527	32.757	69.378	1.00 28.54	AAAA
			LEU A			23.350	33.837	70.058	1.00 27.05	- AAAA
MOTA	1864									
MOTA	1865	CD2	LEU A	235		23.182	31.408	69.558	1.00 27.21	AAAA
MOTA	1866	С	LEU A	235		20.637	30.799	71.429	1.00 30.99	AAAA
ATOM	1867	Ο.	LEU A	235		21.159	29.697	71.547	1.00 31.65	AAAA
	1868	N	GLU A			20.242	31.514	72.471	1.00 31.88	AAAA
MOTA										AAAA
ATOM	1869	CA	GLU A			20.409	31.042	73.838	1.00 32.99	
MOTA	1870	CB	GLU A	236		19.689	31.990	74.790	1.00 34.63	AAAA
ATOM	1871	CG	GLU A	236		19.980	33.449	74.531	1.00 36.79	AAAA
	1872	CD	GLU A			19.044	34.360	75.294	1.00 38.99	AAAA
ATOM							34.303	75.070	1.00 39.03	AAAA
ATOM	1873		GLU A			17.803				
MOTA	1874	OE2	GLU A	236		19.559	35.132	76.126	1.00 41.56	AAAA
ATOM	1875	С	GLU A	236		19.806	29.656	73.982	1.00 32.94	AAAA
ATOM	1876	0	GLU A	236		20.379	28.753	74.595	1.00 31.76	AAAA
		N	LYS A			18.631	29.503	73.399	1.00 32.83	AAAA
MOTA	1877									AAAA
MOTA	1878	CA	LYS A			17.906	28.256	73.471	1.00 33.59	
MOTA	1879	CB	LYS A	237		16.504	28.506	72.942	1.00 35.00	AAAA
ATOM	1880	CG	LYS A	237		15.516	27.436	73.213	1.00 36.69	AAAA
	1881	CD	LYS A			14.310	28.008	73.940	1.00 38.53	AAAA
ATOM							28.331	75.392	1.00 39.27	AAAA
MOTA	1882	CE	LYS A			14.636				
ATOM	1883	NZ	LYS A	237		13.398	28.531	76.204	1.00 39.42	AAAA
MOTA	1884	С	LYS A	237		18.619	27.129	72.707	1.00 33.14	AAAA
ATOM	1885	0	LYS A	237		18.850	26.051	73.260	1.00 33.29	AAAA
	1886	N	SER A			18.985	27.374	71.452	1.00 32.13	AAAA
MOTA								70.685	1,00 31.25	AAAA
MOTA	1887	CA	SER A			19.671	26.345			
MOTA	1888	CB	SER A	238		19.740	26.717	69.194	1.00 30.52	AAAA
MOTA	1889	OG	SER A	238		20.544	27.851	68.970	1.00 29.95	AAAA
	1890	C	SER A			21.075	26.064	71.236	1.00 31.21	AAAA
MOTA						21.556	24.929	71.169	1.00 30.06	AAAA
MOTA	1891	0	SER A							
MOTA	1892	N	LEU A	239		21.740	27.077	71.782	1.00 31.71	AAAA
ATOM	1893	CA	LEU A	239		23.070	26.842	72.351	1.00 33.47	AAAA
ATOM	1894	CB	LEU A	239		23.698	28.13C	72.900	1.00 31.25	AAAA
		CG	LEU A			23.988	29.301	71.977	1.00 29.80	AAAA
ATOM	1895								1.00 29.05	AAAA
MOTA	1896		LEU A			24.589	30.414	72.787		
ATOM	1897	CD2	LEU A	239		24.919	28.903	70.872	1.00 29.36	AAAA
ATOM	1898	С	LEU A	239		22.933	25.839	73.502	1.00 35.41	AAAA
	1899	ō	LEU A			23.812	25.012	73.735	1.00 36.25	AAAA
ATOM							25.906	74.213	1.00 37.34	AAAA
ATOM	1900	N	GLU A			21.816			1.00 37.34	
ATOM	1901	CA	GLU A	240		21.594	25.005	75.331	1.00 39.39	AAAA
ATOM	1902	CB	GLU A			20.281	25.361	76.017	1.00 41.90	AAAA
	1903	CG	GLU A			20.040	24.610	77.308	1.00 45.52	AAAA
MOTA							25.552	78.432	1.00 47.80	AAAA
MOTA	1904	CD	GLU A			19.665				
ATOM	1905	OE1	GLU A	240		18.670	26.295	78.274	1.00 49.70	AAAA
ATOM	1906	OE2	GLU A	240		20.364	25.559	79.469	1.00 48.04	AAAA.
	1907	c	GLU A			21.583	23.555	74.875	1.00 38.80	AAAA
MOTA								75.478	1.00 37.85	AAAA
MOTA	1908	0	GLU A			22.224	22.700			
ATOM	1909	N	ILE A	241		20.847	23.293	73.804	1.00 39.66	AAAA
ATOM	1910	CA	ILE A	241		20.751	21.955	73.223	1.00 40.81	AAAA
	1911	CB	ILE A			19.912	21.994	71.917	1.00 41.10	AAAA
ATOM							20.621	71.287	1.00 40.88	AAAA
MOTA	1912		ILE A			19.850				
MOTA	1913		ILE A			18.502	22.514	72.220	1.00 41.45	AAAA
MOTA	1914	CD1	ILE A	241		17.641	22.745	70.992	1.00 41.14	AAAA

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				•				1 00 41 66	AAAA
ATOM	1915	C :	ILE A 241		22.159	21.424	72.893	1.00 41.66	
	1916	ō ·	ILE A 241	:	22.445	20.229	73.045	1.00 42.10	AAAA
MOTA			VAL A 242		23.026	22.324	72.432	1.00 41.42	AAAA
ATOM	1917				24.394	21.977	72.076	1.00 41.23	AAAA
MOTA	1918		VAL A 242		25.089	23.146	71.351	1.00 40.40	AAAA
MOTA	1919		VAL A 242			22.850	71.171	1.00 39.25	AAAA
MOTA	1920	CG1	VAL A 242		26.556			1.00 39.79	AAAA
ATOM	1921	CG2	VAL A 242		24.438	23.384	70.004		AAAA
ATOM	1922	С	VAL A 242		25.228	21.604	73.298	1.00 42.67	
	1923		VAL A 242		25.882	20.562	73.316	1.00 41.83	AAAA
ATOM			LYS A 243		25.198	22.456	74.318	1.00 44.41	AAAA
MOTA	1924				25.972	22.215	75.523	1.00 46.51	AAAA
ATOM	1925		LYS A 243			23.363	76.522	1.00 47.29	AAAA
MOTA	1926		LYS A 243		25.797		77.664	1.00 48.40	AAAA ·
MOTA	1927	CG	LYS A 243		26.820	23.312		1.00 48.88	AAAA
ATOM	1928	CD	LYS A 243		26.479	24.248	78.823		AAAA
ATOM	1929	CE	LYS A 243		26.355	25.691	78.380	1.00 49.62	AAAA
	1930	NZ	LYS A 243		25.926	26.576	79.505	1.00 50.11	
ATOM		C	LYS A 243		25.639	20.891	76.209	1.00 47.59	AAAA
ATOM	1931		LYS A 243		26.537	20.216	76.711	1.00 48.17	AAAA
MOTA	1932	0			24.362	20.517	76.237	1.00 48.86	AAAA
MOTA	1933	N	GLU A 244			19.262	76.877	1.00 50.82	AAAA
MOTA	1934	ÇA	GLU A 244		23.957		77.103	1.00 52.08	AAAA
MOTA	1935	CB	GLU A 244		22.432	19.208		1.00 53.82	AAAA
ATOM	1936	CG	GLU A 244		21.818	20.405	77.829		
	1937	CD	GLU A 244		20.359	20.174	78.230	1.00 54.49	AAAA
MOTA	1938		GLU A 244		19.666	21.158	78.595	1.00 55.15	AAAA
ATOM			GLU A 244		19.912	19.006	78.200	1.00 54.98	AAAA
MOTA	1939				24.338	18.046	76.033	1.00 51.06	AAAA
MOTA	1940	Ç	GLU A 244		24.206	16.905	76.477	1.00 51.68	AAAA
ATOM	1941	0	GLU A 244				74.820	1.00 51.12	AAAA
ATOM	1942	N	VAL A 245	•	24.810	18.292	73.904	1.00 50.08	AAAA
ATOM	1943	CA	VAL A 245		25.149	17.212	_	1.00 50.22	AAAA
ATOM	1944	CB	VAL A 245		24.217	17.263	72.677		AAAA
	1945		VAL A 245		24.615	16.217	71.651	1.00 51.07	
MOTA	1946		VAL A 245		22.794	17.049	73.118	1.00 50.79	AAAA
ATOM		-	VAL A 245		26.578	17.254	73.397	1.00 49.43	AAAA
ATOM	1947	C	VAL A 245		27.101	16.250	72.917	1.00 48.65	AAAA
MOTA	1948	0			27.220	18.408	73.522	1.00 48.65	AAAA
ATOM	1949	N	PHE A 246			18.552	72.982	1.00 47.97	AAAA
MOTA	1950	CA	PHE A 246		28.556		71.607	1.00 46.45	AAAA
MOTA	1951	CB	PHE A 246		28.420	19.212		1.00 45.35	AAAA
MOTA	1952	CG	PHE A 246		29.553	18.932	70.671	1.00 44.13	AAAA
ATOM	1953	CD1	PHE A 246		29.841	17.629	70.280		AAAA
	1954	CD2			30.291	19.972	70.124	1.00 44.40	
MOTA		CE1			30.840	17.370	69.356	1.00 43.95	AAAA
MOTA	1955	CE2			31.292	19.721	69.197	1.00 43.47	AAAA
MOTA	1956		PHE A 246		31.566	18.422	68.811	1.00 44.05	AAAA
ATOM	1957	CZ			29.481	19.383	73.860	1.00 48.60	AAAA
MOTA	1958	С	PHE A 246				74.239	1.00 49.59	AAAA
ATOM	1959	0	PHE A 246		29.132	20.501	74.198	1.00 48.69	AAAA
MOTA	1960	N	GLU A 247		30.647	18.834		1.00 49.45	AAAA
ATOM.		CA	GLU A 247		31.644	19.578	74.977		AAAA
	1962	CB	GLU A 247		32.174	18.768	76.178		
ATOM	1963	CG	GLU A 247		31.257	18.659	77.398		AAAA
MOTA			GLU A 247		29.986	17.845		1.00 57.34	AAAA
MOTA	1964	CD			29.100	18.315		1.00 58.48	AAAA
MOTA	1965	OE1	GLU A 247		29.877	16.725	77.702		AAAA
MOTA	1966	OE2	GLU A 247				74.024		AAAA
MOTA	1967	С	GLU A 247	'	32.807	19.903		1.00 46.65	AAAA
MOTA	1968	0	GLU A 247	,	33.742				AAAA
MOTA	1969		PRO A 248		32.748		73.371		
	1970		PRO A 248		31.651	22.033	73.543		AAAA
ATOM			PRO A 248		33.710		72.411		AAAA
MOTA	1971				33.063			1.00 45.57	AAAA
MOTA	1972		PRO A 248						AAAA
ATOM	1973		PRO A 248		31.604		·		AAAA
ATOM	1974	C	PRO A 248		35.155				AAAA
ATOM	1975	_	PRO A 24	3	35.401			T.00 44.31	AAAA
	1976		GLU A 24		36.100			1.00 42.21	
ATOM			GLU A 24		37.522				AAAA
ATOM	1977		GLU A 24		38.344			1.00 39.58	AAAA
MOTA	1978				37.960			7 1.00 41.32	AAAA
MOTA	1979		GLU A 24						AAAA
ATOM	1980	CD	GLU A 24	,	38.825	19.00	, , , , , , , , ,		
				·					
=			TERILE	TITE	CHEL	r form i	761		

MOTA	1981	OE1	GLU A	249	38.871	18.017	69.996	1.00 41.55	AAAA
MOTA	1982	OE2	GLU A	249	39.462	17.187	71.926	1.00 42.49	AAAA
ATOM	1983	С	GLU A	249	37.840	22.873	71.718	1.00 38.81	AAAA
MOTA	1984	0	GLU A	249	38.715	23.617	72.181	1.00 38.27	AAAA
ATOM	1985	N	VAL A	250	37.109	23.160	70.644	1.00 36.60	AAAA
MOTA	1986	CA	VAL A		37.242	24.402	69.890	1.00 34.20	AAAA
ATOM	1987	CB	VAL A		38.379	24.321	68.862	1.00 33.73	AAAA
MOTA	1988		VAL A		38.085	23.209	67.864	1.00 33.26	AAAA
	1989		VAL A.		38.546	25.678	68.153	1.00 33.75	AAAA
MOTA	1990	C	VAL A		35.945	24.617	69.130	1.00 31.98	AAAA
MOTA		ō	VAL A		35.205	23.658	68.904	1.00 32.36	AAAA
MOTA	1991 1992		TYR A		35.657	25.863	68.760	1.00 28.65	AAAA
ATOM		N CA	TYR A		34.449	26.150	67.991	1.00 26.49	AAAA
MOTA	1993		TYR A		33.241	26.442	68.906	1.00 24.32	AAAA
MOTA	1994	CB	TYR A		33.193	27.853	69.465	1.00 22.96	AAAA
MOTA	1995	CG			32.771	28.931	68.668	1.00 22.21 :	
ATOM	1996		TYR A			30.234	69.151	1.00 21.29	AAAA
MOTA	1997		TYR A		32.791		70.771	1.00 21.47	AAAA
MOTA	1998		TYR A		33.628	28.124	71.265	1.00 20.80	AAAA
MOTA	1999	CE2	TYR A		33.651	29.425		1.00 20.77	AAAA
MOTA	2000	CZ	TYR A		33.237	30.475	70.449	1.00 20.77	AAAA
MOTA	2001	OH	TYR A		33.309	31.768	70.913	1.00 24.59	AAAA
ATOM	2002	С	TYR A		34.691	27.345	67.092		
MOTA	2003	0	TYR A		35.504	28.216	67.410		AAAA
MOTA	2004	N	LEU A		33.984	27.374	65.970	1.00 22.49	AAAA
MOTA	2005	CA	LEU A		34.082	28.482	65.045	1.00 20.96	AAAA
MOTA	2006	CB	LEU A		34.523		63.657	1.00 21.31	AAAA
MOTA	2007	CG	LEU A		35.940	27.472	63.556	1.00 21.03	AAAA
MOTA	2008		LEU A		35.947	26.028	63.977	1.00 22.16	AAAA
MOTA	2009	CD2	LEU A		36.440	27.594	62.143	1.00 22.13	AAAA
MOTA	2010	C	LEU A		32.731	29.159	64.959	1.00 19.60	AAAA
MOTA	2011	0	LEU A	252	31.689	28.523		1.00 19.95	AAAA
MOTA	2012	N	LEU A	253	32.748	30.461	64.756	1.00 17.95	AAAA
MOTA	2013	CA	LEU A		31.521	31.222	64.675	1.00 17.33	AAAA
ATOM	2014	CB	LEU A	253	31.441	32.142	65.900	1.00 16.31	AAAA
MOTA	2015	CG	LEU A		30.266	33.070	66.153	1.00 15.81	AAAA
ATOM	2016	CD1	LEU A	253	28.990	32.267	66.377	1.00 14.74	AAAA
ATOM	2017	CD2	LEU A	253	30.602	33.925	67.368	1.00 15.83	AAAA
MOTA	2018	С	LEU A	253	31.564	32.035	63.386	1.00 16.60	AAAA
MOTA	2019	0	LEU A	253	32.548	32.722	63.132	1.00 16.40	AAAA
MOTA	2020	N	GLN A		30.526	31.936	62.557	1.00 15.88	AAAA
MOTA	2021	CA	GLN A		30.507	32.716	61.328	1.00 16.27	AAAA
ATOM	2022	CB	GLN A		30.045	31.881	60.121	1.00 15.88	AAAA
MOTA	2023	CG	GLN A	254	28.587	32.048	59.734	1.00 18.52	AAAA
MOTA	2024	CD	GLN A	254	28.380	32.935	58.519	1.00 17.54	AAAA
ATOM	2025	OE1	GLN A	254	28.714	32.572	57.391	1.00 15.89	AAAA
ATOM	2026	NE2	GLN A	254	27.828	34.103	58.750		AAAA
ATOM	2027	С	GLN A	254	29.527	33.825	61.650	1 00 16.91	AAAA
MOTA	2028	0	GLN A	254	28.450	33.571	62.198	1 00 17.41	AAAA
ATOM	2029	N	LEU A	255	29.911	35.053	61.319	1.00 16.68	AAAA
MOTA	2030	CA	LEU A	255	29.102	36.215	61.619	1.00 16.42	AAAA
MOTA	2031	CB	LEU A	255	29.861	37.080	62.616	1.00 14.93	AAAA
MOTA	2032	·CG	LEU A	255	30.269	36.301	63.860	1.00 13.90	AAAA
MOTA	2033	CD1	LEU A	255	31.494	36.924	64.515	1.00 12.24	AAAA
ATOM	2034	CD2	LEU A	255	29.083	36.202	64.774	1.00 12.80	AAAA
ATOM	2035	C	LEU A		28.699	37.048	60.404	1.00 18.32	AAAA
ATOM	2036	ō	LEU A		29.170	38.177	60.216	1.00 17.59	AAAA
	2037	N	GLY A		27.813	36.482	59.588	1.00 19.75	AAAA
MOTA	2038	CA	GLY A	256	27.322	37.188	58.422	1.00 20.77	AAAA
MOTA	2039	c	GLY A	256	26.422	38.302	58.927	1.00 21.73	AAAA
MOTA	2040	ō	GLY A	256	25.642	38.096	59.857	1.00 21.38	AAAA
MOTA	2041	N	THR A	257	26.528	39.485	58.325	1.00 22.82	AAAA
MOTA	2041	CA	THR A		25.721	40.622	58.746	1.00 23.85	AAAA
MOTA	2042	CB	THR A		26.460	41.968	58.549	1.00 23.99	AAAA
MOTA		061	THR A	257	26.729	42.169	57.153	1.00 25.54	AAAA
ATOM	2044		THR A		27.780	41.985	59.329	1.00 24.07	AAAA
MOTA	2045 2046	C GZ			24.438	40.691	57.948	1.00 24.97	AAAA
MOTA	2090	٠.			, •	/	•		•

MOTA	2047	0	THR	Α	257		23.692	41.672	58.048	1.00 25.84	AAAA
ATOM	2048	N	ASP				24.152	39.665	57.154	1.00 25.18	AAAA
ATOM	2049	CA	ASP				22.935	39.753	56.379	1.00 26.18	AAAA
ATOM	2050	CB	ASP				22.950	38.830	55.149	1.00 25.52	AAAA
	2051	CG	ASP				23.211	37.392	55.494	1.00 26.33	AAAA
ATOM	2052	C			258		21.649	39:574	57.178	1.00 26.74	AAAA
MOTA		0			258		20.571	39.823	56.643	1.00 26.57	AAAA
ATOM	2053.		ASP				23.014	37.029	56.675	1.00 26.85	AAAA
MOTA	2054						23.585	36.623	54.572	1.00 24.06	AAAA
MOTA	2055		ASP				23.363	39.114	58.449	1.00 26.95	AAAA
MOTA	2056	N			259		22.834	38.589	59.271	1.00 27.34	AAAA
MOTA	2057	CD			259	-			59.190	1.00 27.05	AAAA
MOTA	2058	CA			259		20.467	38.976	60.425	1.00 26.38	AAAA
MOTA	2059	CB			259		20.886	38.186	60.669	1.00 27.84	AAAA
ATOM	2060	CG			259		22.247	38.718	59.543	1.00 27.32	AAAA
ATOM	2061	C			259		19.914	40.365	59.871-	1.00 27.32	AAAA
ATOM	2062	0			259		18.739	40.510	-	1.00 26.97	AAAA
ATOM	2063	N			260		20.771	41.383	59.452	1.00 26.74	AAAA
MOTA	2064	CA			260		20.389	42.763	59.752	1.00 27.21	AAAA
MOTA	2065	CB			260		21.621	43.680	59.680		
MOTA	2066	CG			260		22.732	43.465	60.709	1.00 27.01	AAAA
MOTA	2067		LEU				23.889	44.380	60.408	1.00 25.51	AAAA
ATOM	2068		LEU				22.189	43.718	62.112	1.00 27.39	AAAA
MOTA	2069	C			260		19.295	43.351	58.865	1.00 26.47	AAAA
ATOM	2070	0			260		19.278	43.137	57.649	1.00 26.72	AAAA
MOTA	. 2071	N			261		18.413	44.126	59.494	1.00 26.32	AAAA
MOTA	2072	CA			261		17.283	44.808		1.00 27.20	AAAA
ATOM	2073	CB			261		16.732	45.885	59.780	1.00 28.71	AAAA
MOTA	2074	CG			261		15.644	46.789	59.190	1.00 29.24	AAAA
ATOM	2075		LEU				14.433	45.954	58.883	1.00 29.44	AAAA
ATOM -	2076	CD2	LEU				15.284	47.906	60.162	1.00 29.72	AAAA
MOTA	2077	С			261		17.506	45.454	57.473	1.00 27.90	AAAA
ATOM	2078	0			261		16.675	45.294	56.577	1.00 28.21	AAAA
ATOM	2079	N			262			46.202	57.310	1.00 27.61	AAAA AAAA
ATOM	2080	CA			262	•	18.887	46.877	56.043	1.00 26.92 1.00 25.85	AAAA
ATOM	2081	CB			262.		19.949	47.955	56.241 57.117	1.00 25.36	AAAA
MOTA	2082	CG			262		19.549	49.119	58.580	1.00 25.78	AAAA
MOTA	2083	CD			262		19.552	48.787 47.631	58.938	1.00 24.64	AAAA
ATOM	2084		GLU				19.859 19.255	49.694	59.381	1.00 25.82	AAAA
MOTA	2085	-	GLU				19.346	45.995	54.882	1.00 28.79	AAAA
MOTA	2086	C			262		19.354	46.439	53.724	1.00 28.70	AAAA
MOTA	2087	0			262 263		19.743	44.758	55.179	1.00 29.57	AAAA
MOTA	2088	N			263		20.230	43.853	54.145	1.00 28.99	AAAA
MOTA	2089 2090	CA CB			263		21.160	42.802	54.760	1.00 27.89	AAAA
MOTA	2091	CG			263		21.986	42.062	53.714	1.00 29.02	AAAA
MOTA	2091				263		23.194	41.863	53.957	1.00 28.06	AAAA
ATOM	2093				263		21.438	41.663	52.660	1.00 28.80	AAAA
ATOM	2094	Č			263		19.066	43.197	53.431	1.00 29.73	. AAAA
MOTA	2095	ŏ			263		18.258	42.510	54.043	1.00 29.15	AAAA
MOTA	2096	N			264		19.002	43.416	52.122	1.00 31.25	AAAA
MOTA	2097	CA			264		17.925	42.888	51.306	1.00 32.43	AAAA
ATOM	2098	CB	TVD	Λ	264		17.913	43.558	49.938	1.00 34.53	AAAA
MOTA	2099	CG			264		17.627	45.038	49.997	1.00 38.21	AAAA
MOTA	2100		TYR				18.664		49.983	1.00 39.87	AAAA
ATOM	2101		TYR				18.409	47.335	50.068	1.00 41.74	AAAA
ATOM		CD2			264		16.316	45.511	50.103	1.00 40.10	AAAA
ATOM	2102	CES			264		16.044	46.877	50.191	1.00 41.50	AAAA
MOTA	2103				264		17.095	47.786	50.170	1.00 42.75	AAAA
MOTA	2104	CZ			264		16.838	49.147		1.00 44.65	AAAA
MOTA	2105	он					17.897	41.385	51.135	1.00 32.50	AAAA
ATOM	2106	C			264		16.819	40.816	50.968	1.00 32.49	AAAA
ATOM	2107	0			264		19.064	40.740		1.00 32.64	AAAA
ATOM	2108	И	- TEO	A	265		19.122	39.281		1.00 31.92	AAAA
MOTA	2109	CA			265		20.525	38.823	• _	1.00 32.75	AAAA
ATOM	2110	CB			265		20.323	39.010			AAAA
ATOM	2111	CG			265		22.213	38.588		1.00 31.59	AAAA
MOTA	2112	CDI	. LEU		203		66.613	50.500			•

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Figure 19-33

ATOM	2113	CD2	LEU A	265	19.803	38.166	48.361	1.00 34.62	AAAA
									AAAA
MOTA	2114	С	LEU A	265	18.693	38.540	52.296	1.00 30.33	
MOTA	2115	0	LEU A	265	19.024	37.375	52.484	1.00 30.30	AAAA
	2116	N	SER A		17.945	39.230	53.147	1.00 29.23	AAAA
MOTA									
MOTA	2117	CA	SER A	266	17.434	38.649	54.371	1.00 29.72	AAAA
ATOM -	2118	CB	SER A	266	18.398	38.894	55.519	1.00 32.09	AAAA
		ŌĠ	SER A		17.728	38.810	56.771	1.00 33.43	AAAA
MOTA	2119								
ATOM	2120	C	SER A	266	16.115	39.290	54.698	1.00 28.71	ÀAAA
ATOM	2121	0	SER A	266	15.924	40.473	54.444	1.00 29.67	AAAA
							55.276	1.00 27.82	AAAA
ATOM	2122	N	LYS A	267	15.209	38.517			
ATOM	2123	CA	LYS A	267	13.908	39.045	55.654	1.00 27.56	AAAA
	2124	CB	LYS A	267	12.821	38.076	55.222	1.00 28.75	AAAA
MOTA									
MOTA	2125	CG	LYS A		12.733	37.922		1.00 29.67	AAAA
MOTA	2126	CD	LYS A	267	12.343	39.223	53.053	1.00 30.13	AAAA
		CE	LYS A		12.303	39.036	51.546	1.00 31.86	AAAA
MOTA	2127								
ATOM	2128	NZ	LYS A	267	11.796	40.252	50.843	1.00 33.92	- AAAA
ATOM	2129	С	LYS A	267	13.800	39.327	57.152	1.00 27.18	· AAAA
			LYS A		12.707	39.591	57.665	1.00 27.18	AAAA
MOTA	2130	0							
MOTA	2131	N	PHE A	268	14.944	39.267	57.836	1.00 26.12	AAAA
MOTA	2132	CA	PHE A	268	15.048	39.532	59.271	1.00 25.72	AAAA
			PHE A		16.272	38.830	59.856	1.00 24.94	AAAA
MOTA	2133	CB							
MOTA	2134	CG	PHE A	268	16.167	37.334	59.896	1.00 25.07	AAAA
MOTA	2135	CD1	PHE A	268	17.271	36.565	60.267	1.00 24.56	AAAA
					14.955		59.629	1.00 23.76	AAAA
ATOM	2136	CD2	PHE A			36.687			
MOTA	2137	CE1	PHE A	268	17.174	35.169	60.384	1.00 23.71	AAAA
ATOM	2138	CE2	PHE A	268	14.850	35.303	59.739	1.00 23.86	AAAA
						34.542	60.121	1.00 23.68	AAAA
MOTA	2139	cz	PHE A		15.966				
ATOM	2140	С	PHE A	268	15.190	41.030	59.513	1.00 25.77	AAAA
MOTA	2141	0	PHE A	268	15.811	41.734	58.726	1.00 25.81	AAAA
						41.524	60.595	1.00 26.02	AAAA
MOTA	2142	N	ASN A						
MOTA	2143	CA	ASN A	269	14.718	42.943	60.890	1.00 26.58	AAAA
MOTA	2144	CB	ASN A	269	13.330	43.584	61.058	1.00 25.47	AAAA
						43.252.	59.906	1.00 25.37	AAAA
MOTA	2145	CG	ASN A		12.379				
MOTA	2146	OD1	ASN A	269	12.761	43.260	58.734	1.00 23.82	AAAA
ATOM	2147	MD2	ASN A	269	11.123	42.985	60.245	1.00 24.03	AAAA
					15.540	43.112	62.169	1.00 26.82	AAAA
MOTA	2148	С	ASN A						
MOTA	2149	0	ASN A	269	15.089	43.715	63.150	1.00 27.98	AAAA
MOTA	2150	N	LEU A	270	16.744	42.559	62.149	1.00 26.07	AAAA
					17.639	42.642	63.289	1.00 25.97	AAAA
MOTA	2151	CA	LEU A						
ATOM	2152	CB	LEU A	270	18.634	41.479	63.265	1.00 23.76	AAAA
MOTA	2153	CG	LEU A	270	18.048	40.070	63.225	1.00 23.36	AAAA
			LEU A		19.115	39.090	63.710	1.00 21.90	AAAA
MOTA	2154								
ATOM	2155	CD2	LEU A	270	16.824	39.971	64.122	1.00 22.05	AAAA
ATOM	2156	С	LEU A	270	18.420	43.961	63.360	1.00 27.13	AAAA
-			LEU A			44.750	62.399	1.00 25.99	AAAA
MOTA	2157	0			18.475				
ATOM	2158	N	SER A	271	19.038	44.176	64.517	1.00 27.97	. AAAA
ATOM	2159	CA	SER A	271	19.832	45.370	64.767	1.00 27.95	AAAA
			SER A		19.235	46.137	65.943	1.00 27.32	AAAA
MOTA	2160	CB							
ATOM	2161	OG	SER A	271	19.184	45.297	67.089	1.00 27.90	AAAA
MOTA	2162	С	SER A	271	21.276	44.987	65.084	1.00 28.15	AAAA
	2163	ŏ	SER A		21.574	43.832	65.401	1.00 26.99	AAAA
MOTA									AAAA
MOTA	2164	N	ASN A	272	22.156	45.980	64.979	1.00 28.96	
MOTA	2165	CA	ASN A	272	23.590	45.861	65.266	1.00 29.54	AAAA
					24.247	47.243	65.223	1.00 30.96	AAAA
ATOM	2166	CB	ASN A						
MOTA	2167	CG	ASN A		24.647	47.640	63.850	1.00 31.20	AAAA
	2168	OD1	ASN A		24.960	48.794	63.594	1.00 31.73	AAAA
ATOM		1750	3 027 -	272	_	46.674	62.948	1.00 31.93	AAAA
ATOM	2169		ASN A		24.670				
MOTA	2170	С	ASN A	272	23.821	45.309	66.645	1.00, 29.63	AAAA
	2171	0	ASN A		24.574	44.361	66.843	1.00 29.85	AAAA
MOTA							67.600	1.00 29.77	AAAA
MOTA	2172	N	VAL A		23.180	45.959			
ATOM	2173	CA	VAL A	273	23.290	45.602	68.994	1.00 30.89	AAAA
	2174	CB	VAL A		22.436	46.576	69.816	1.00 31.61	AAAA
MOTA							71.293	1.00 33.17	AAAA
MOTA	2175		VAL A		22.716	46.403			
ATOM	2176	CG2	VAL A	273	22.740	47.998	69.372	1.00 31.82	AAAA
	2177	c	VAL A		22.883	44.144	69.266	1.00 30.74	AAAA
MOTA									AAAA .
ATOM	2178	0	VAL A	273	23.550	43.431	70.022	1.00 31.23	· MANA

	2170	NT	ALA.	a 2	74	21.785	4:	3.706	68.659	1.00	30.25	AAAA
MOTA	2179 2180	N CA	ALA .			21.327		2.333	68.840		29.87	AAAA
MOTA	2180	CB	ALA.			20.005		2.119	68.112		29,64	AAAA
MOTA	2182	C	ALA .			22.395	4	1.438	68.247		29.35	AAAA
ATOM	2182	ō	ALA .			22.707	4	0.373	68.778		29.18	AAAA
MOTA	2184	И	PHE	A 2	75	22.946	4	1.893	67.127	1.00	29.30	AAAA
MOTA	2185	CA	PHE			23.991	4	1.170	66.428	1.00	28.91	AAAA
ATOM	2186	CB	PHE	A 2	275	24.375		1.909	65.150		28.77	AAAA
MOTA	2187	CG	PHE			25.354	4	1.170	64.308		28.08	AAAA
MOTA	2188	CD1	PHE			25.015		9.954	63.740		28.92	AAAA
MOTA	2189	CD2	PHE	A 2	275	26.621	4	1.684	64.077		29.48	AAAA
MOTA MOTA	2190	CE1	PHE	A 2	275	25.928	. 3	9.259	62.945		29.20	AAAA
	2191		PHE			27.546	4	0.988	63.279		29.24	AAAA
MOTA MOTA	2192	CZ	PHE			27.193	- 3	9.779	62.716		28.30	AAAA
ATOM	2193	c	PHE	A :	275	25.196	4	1.058	67.351		27.64	AAAA
ATOM	2194	ō	PHE			25.728		9.975	67.558		28.65	AAAA
MOTA	2195	N	LEU	A :	276	25.606		2.189	67.902		26.81	AAAA AAAA
ATOM	2196	CA	LEU	A :	276	26.732		2.260	68.831		27.38	AAAA AAAA
MOTA	2197	CB	LEU	Α :	276	26.878		3.700	69.353	1.00	27.53 26.37	AAAA
ATOM	2198	CG	LEU	A :	276	28.202		4.213	69.928			AAAA
ATOM	2199	CD1	LEU	A	276	27.923		5.488	70.721	1.00	25.71 27.06	AAAA
ATOM	2200	CD2	LEU	Α.	276	28.842		3.189	70.827	1.00	26.49	AAAA
ATOM	2201	C	LEU	A :	276	26.486		1.317	70.021		25.26	AAAA
ATOM	2202	0	LEU	Α	276	27.387		10.603	70.471		27.46	AAAA
ATOM	2203	N	LYS			25.257		11.322	70.524		28.63	AAAA
MOTA	2204	CA	LYS			24.894		10.468	71.642		30.63	AAAA
MOTA	2205	CB	LYS	Α	277	23.542		10.862	72.223 73.153		33.14	AAAA
ATOM	2206	CG	LYS	A	277	23.590		12.029	74.268		34.94	AAAA
ATOM	2207	CD	LYS			22.599		11.791	75.029		36.17	AAAA
ATOM	2208	CE	LYS			22.964		40.519	76.104		0 38.64	AAAA
ATOM	2209	NZ	LYS	A	277	21.979	-	38.997	71.297		0 28.53	AAAA
ATOM	2210	С	LYS			24.846		38.152	72.146		0 28.45	AAAA
MOTA	2211	0	LYS			25.118 24.46		38.681	70.064		0 28.47	AAAA
ATOM	2212	N	ALA			24.404		37.280	69.656		0 27.66	AAAA
MOTA	2213	CA	ALA			23.94		37.181	68.201		0 26.40	AAAA
MOTA	2214	CB	ALA ALA			25.83		36.754	69.820		0 26.63	AAAA
MOTA	2215	C	ALA			26.08	-	35.644	70.317		0 25.19	AAAA
MOTA	2216	N	PHE			26.76		37.616	69.427		0 26.50	AAAA
MOTA	2217 2218	CA	PHE			28.18	1	37.345	69.481	1.0	0 25.83	AAAA
MOTA	2219	CB	PHE			28.93	4	38.521	68.869		0 26.35	AAAA
MOTA	2220	CG			279	30.41	3	38.319	68.796		0 27.92	AAAA
MOTA	2221	CD	PHE	A	279	30.94		37.256		1.0	0 28.58	AAAA
ATOM ATOM	2222	CD2	PHE	A	279	31.28		39.201		1.0	0 28.33	AAAA
MOTA	2223	CE:				32.33		37.078			0 28.22	AAAA
ATOM	2224	CE		Α	279	32.66		39.030			0 28.11	AAAA AAAA
MOTA	2225	CZ			279	33.18	5	37.968		_	0 28.21 0 25.47	
ATOM	2226	С	PHE	Α	279	28.66		37.118			0 24.32	AAAA
MOTA	2227	0	PHE	Α	279	29.28		36.091			0 25.12	AAAA
ATOM	2228	N	ASN	Α	280	28.38		38.075			0 25.05	AAAA
ATOM	- 2229	CA			280	28.84		37.944			0 24.42	AAAA
ATOM	2230	CB	ASN	Α	280	28.70		39.269		1 (0 24.56	AAAA
ATOM	2231	CG	ASN	Α	280	29.68		40.300		1 (0 23.24	AAAA
MOTA	2232	OD:	l ASN	A	280	30.84		39.980 41.543			0 24.59	AAAA
ATOM	2233	ND:	2 ASN	Α	280	29.23		36.814		1.0	0 24.79	AAAA
ATCM	2234	С	ASN	A	280	28.21		36.272			0 24.96	AAAA
MOTA	2235	0			280	28.82		36.444			00 24.87	AAAA
ATOM	2236				281	26.99 26.33		35.337		1.0	00 24.80	AAAA
ATOM	2237				281		_	35.252			00 24.40	AAAA
ATOM	2238		ILE	A	281	24.86		33.907		1 1.0	00 25.03	AAAA
MOTA	2239		2 ILE	A	201 201	24.23		36.386		1.0	00 24.70	AAAA
ATOM	2240		1 ILE	. A	201	22.63		36.379		1.	00 26.49	AAAA
MOTA	2241	_	1 ILE	. A	201	27.04		34.02		1 1.	00 25.21	AAAA
MOTA	2242				281	27.22		33.170		1.	00 24.97	AAAA
MOTA	2243				281 282	27.44		33.866			00 25.98	
ATCM	2244	N	VAL		202	27.35			-			•
	•		•									

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Figure 19-35

					•				
MOTA	2245	CA	VAL A	A 282	28.150	32.656	72.193	1.00 25.15	AAAA
ATOM	2246	CB	VAL A	A 282	28.451	32.666	70.677	1.00 23.83	AAAA
MOTA	2247	CG1	VAL A	A 282	29.315	31.470	70.311	1.00 23.58	AAAA
MOTA	2248	CG2	VAL A		27.173	32.633	69.899	1.00 22.73	AAAA
MOTA	2249	С		A 282	29.478	32.553	72.936	1.00 25.73	AAAA
MOTA	2250	0		A 282	29.928	31.457	73.275	1.00 25.31	AAAA
MOTA	2251	N	ARG A		30.100	33.702	73.176	1.00 26.90 1.00 28.87	AAAA AAAA
MOTA	2252	CA		A 283	31.372	33.760	73.885 73.684	1.00 28.16	AAAA
MOTA	2253	CB	ARG A		32.027 32.364	35.131 35.440	72.240	1.00 27.22	AAAA '
MOTA	2254	CG		A 283 A 283	32.821	36.862	72.098	1.00 27.08	AAAA
MOTA	2255 2256	CD NE		A 283	34.035	37.116	72.854	1.00 26.73	AAAA
MOTA	2257	CZ		A 283	34.514	38.327	73.091	1.00 26.82	AAAA
MOTA . MOTA	2258	NHI			33.873	39.384	72.626	1.00 27.36	AAAA
ATOM	2259	NH2		A 283	35.622	38.484	73.798	1.00 26.95	AAAA
ATOM	2260	С		A 283	31.183	33.494	75.376	1.00 30.71	AAAA
ATOM	2261	0	ARG I	A 283	32.086	32.981	76.027	1.00 30.68	AAAA
ATOM	2262	N	GLU I	A 284	30.014	33.842	75.911	1.00 32.71	AAAA
ATOM	2263	CA		A 284	29.735	33.623	77.323	1.00 35.53	AAAA
ATOM	2264	CB		A 284	28.482	34.391	77.751	1.00 37.39	AAAA
MOTA	2265	CG		A 284	28.538	35.854	77.392	1.00 41.73	AAAA
MOTA	2266	CD		A 284	27.272	36.631	77.754	1.00 45.27 1.00 46.66	AAAA AAAA
MOTA	2267		GLU I		26.151	36.078 37.817	77.610 78.148	1.00 46.88	AAAA
MOTA	2268		GLU A	A 284 A 284	27.405 29.524	32.133	77.564	1.00 36.25	AAAA
ATOM	2269 2270	0		A 284	29.920	31.593	78.601	1.00 37.85	AAAA
MOTA	2271	N		A 285	28.916	31.464	76.591	1.00 35.24	AAAA
MOTA MOTA	2272	CA			. 28.637	30.041	76.708	1.00 33.88	AAAA
ATOM	2273	CB		A 285	27.505	29.619	75.737	1.00 33.71	AAAA
ATOM	2274		VAL	A 285	27.201	28.137	75.888	1.00 32.59	AAAA
ATOM	2275	CG2	VAL A	A 285	26.254	30.457	76.001	1.00 32.77	AAAA
ATOM	2276	С	VAL	A 285	29.847	29.149	76.456	1.00 33.47	AAAA
MOTA	2277	0		A 285	30.140	28.262	77.257	1.00 34.23	AAAA
ATOM	2278	N		A 286	30.568	29.389	75.364 75.036	1.00 32.34 1.00 29.92	AAAA AAAA
MOTA	2279	CA		A 286	31.706	28.535 27.960	73.635	1.00 29.77	AAAA
ATOM	2280	CB CG		A 286 A 286	31.533 30.267	27.179	73.444	1.00 28.64	AAAA
MOTA	2281 2282		PHE		29.152	27.772	72.863	1.00 28.75	AAAA
ATOM	2283		PHE		30.197	25.837	73.827	1.00 28.55	AAAA
ATOM ATOM	2284	CE1		A 286	27.983	27.039	72.660	1.00 29.04	AAAA
ATOM	2285		PHE		29.037	25.095	73.629	1.00 28.19	AAAA
ATOM	2286	CZ	PHE :	A 286	27.929	25.694	73.045	1.00 28.73	AAAA
ATOM	2287	С		A 286	33.106	29.113	75.132	1.00 29.13	AAAA
MOTA	2288	0		A 286	34.073	28.436	74.760	1.00 28.54	AAAA
MOTA	2289	N		A 287	33.224	30.341	75.637 75.744	1.00 28.42 1.00 27.07	AAAA AAAA
MOTA	2290	CA		A 287	34.525	30.987 31.611	74.419	1.00 26.64	AAAA ·
MOTA	2291	C		A 287 A 287	34.932 34.088	32.042	73.649	1.00 27.13	AAAA
MOTA	2292 2293	И О		A 288	36.227	31.665	74.146	1.00 27.20	AAAA
MOTA	2294	CA		A 288	36.719	32,238	72.900	1.00 27.52	AAAA
MOTA MOTA	2295	CB		A 288	38.073	32.923	73.108	1.00 28.18	AAAA
MOTA	2296	ĊĠ		A 288	38.036	34.177	73.959	1.00 28.88	AAAA
ATOM	2297	CD		A 288	37.329	35.330	73.279	1.00 29.58	AAAA
ATOM	2298		GLU .		37.807	35.813	72.243	1.00 29.94	AAAA
ATOM	2299	OE2	GLU :	A 288	36.281	35.761	73.782	1.00 31.89	AAAA
ATOM	2300	С		A 288	36.877	31.158	71.843	1.00 27.44	AAAA
ATOM .	2301	0		A 288	37.169	30.007	72.162	1.00 27.87	AAAA
MOTA	2302	N		A 289	36.663	31.547	70.589	1.00 26.55	AAAA
MOTA	2303	CA		A 289	36.795	30.638	69.466 68.254	1.00 25.25 1.00 24.55	AAAA AAAA
MOTA	2304	Ç		A 289	37.285	31.414 32.586	68.369	1.00 24.30	AAAA
MOTA	2305	0		A 289	37.635	32.386	67.095	1.00 24.04	AAAA
MOTA	2306	N		A 290 A 290	37.320 37.756	31.407	65.863	1.00 23.76	AAAA
MOTA	2307	CA CB		A 290 A 290	38.288	30.346	64.867	1.00 24.94	AAAA
MOTA	2308 2309		VAL		38.835	31.012	63.596	1.00 22.73	AAAA
MOTA	2310		VAL .		39.375	29.506	65.555	1.00 24.74	AAAA
МОТА	- J - U						-		•

		-	VAL A	200 '	36.536	32.122	65.277	1.00 23.90	AAAA
MOTA	2311	C	VAL A	290	35.497	31.502	65.100	1.00 25.15	AAAA
ATOM	2312				36.662	33.415	64.976	1.00 23.09	AAAA
MOTA	2313		TYR A		35.544	34.211	64.446	1.00 21.41	AAAA
MOTA	2314				35.472	35.540	65.193	1.00 20.57	AAAA
MOTA	2315	CB	TYR A		35.511	35:346	66.677	1.00 19.87	AAAA
MOTA	2316		TYR A		36.596	35.782	67.432	1.00 20.86	AAAA
ATOM	2317	CD1	TYR A	291	36.677	35.513	68.793	1.00 21.47	AAAA
MOTA	2318	CEI	TYR A	. 291	34.509	34:647	67.318	1.00 20.90	AAAA
MOTA	2319	CD2	TYR A	. 291	34.579	34.372	68.675	1.00 21.90	AAAA
MOTA	2320		TYR A			34.800	69.403	1.00 21.25	AAAA
ATOM	2321	CZ	TYR A		35.661	34.469	70.730	1.00 23.75	AAAA
MOTA	2322	OH	TYR A		35.737	34.483	62.946	1.00 21.25	AAAA
ATOM	2323	C	TYR A		35.607	35.077	62.451	1.00 21.10	AAAA
MOTA	2324	0	TYR A		36.573	34.084	62.231	1.00 20.92	AAAA
MOTA	2325	N	LEU A		34.557	34.260	60.779-	1.00 20.92	AAAA
MOTA	2326	CA	LEU A		34.518	32.916	60.080	1.00 19.93	AAAA
MOTA	2327	CB	LEU A		34.235	31.688	60.399	1.00 17.31	AAAA
ATOM	2328	CG	LEU A		35.104	30.515	59.528	1.00 16.05	AAAA
MOTA	2329	CD1	LEU A	292	34.685	32.000	60.163	1.00 18.07	AAAA
ATOM	2330		LEU A	1 292	36.552	35.288	60.283	1.00 21.12	AAAA
MOTA	2331	С	LEU A	1 292	33.515		61.020	1.00 20.70	AAAA
MOTA	2332	0	LEU A		32.652	35.741	59.017	1.00 21.74	AAAA
ATOM	2333	N	GLY F		33.660	35.660	58.410	1.00 21.48	AAAA
MOTA	2334	CA	GLY 7	1 293	32.752	36.612	57.770	1.00 21.65	AAAA
ATOM	2335	С	GLY A	A 293	31.612		58.235	1.00 22.25	AAAA
MOTA	2336	0	GLY A		31.237	34.790	56.691	1.00 22.66	AAAA
ATOM	2337	N		A 294	31.060			1.00 23.61	AAAA
MOTA	2338	CA	GLY A		29.957		56.034	1.00 24.56	AAAA
ATOM	2339	С	GĻY 1	A 294	29.180		55.146	1.00 25.54	AAAA
ATOM -	2340	0	GLY A	A 294	29.679		54.790	1.00 24.06	AAAA
ATOM	2341	N	GLY A	A 295	27.956		54.794		AAAA
ATOM	2342	CA	GLY A	A 295	27.139		53.927	1.00 22.78	AAAA
ATOM	2343	С	GLY Z	A 295	26.902		54.483	1.00 23.11	AAAA
ATOM	2344	0		A 295	26.870		55.696	1.00 22.87	AAAA
MOTA	2345	N	GLY 2	A 296	26.733	39.442	53.584	1.00 22.78	AAAA
MOTA	2346	CA	GLY 2	A 296	. 26.497		53.993	1.00 23.44	AAAA
ATOM	2347	c	GLY 2	A 296	26.471		52.718	1.00 23.72	AAAA
MOTA	2348	0	GLY .	A 296	27.474			1.00 23.73	AAAA
ATOM	2349	N		A 297	25,356			1.00 23.41	AAAA
ATOM	2350	CA		A 297	25.282	42.991	51.163	1.00 22.71	AAAA
ATOM	2351	CB		A 297	24.252		50.296	1.00 21.55	AAAA
ATOM	2352	CG	TYR .	A 297	24.496		50.317	1.00 21.93	AAAA
ATOM	2353	CD1	TYR	A 297	24.036				AAAA
MOTA	2354	CE1	TYR	A 297	24.400			1.00 21.59	AAAA
MOTA	2355	CD2	TYR	A 297	25.320				AAAA
MOTA	2356	CE2	TYR	A 297	25.688				AAAA
	2357	CZ	TYR	A 297	125.242	38.127	50.511		AAAA
MOTA MOTA	2358	OH	TYR	A 297	25.72	L 36.8 41			
	2359	C	TYR	A 297	25.042				AAAA
MOTA	2360	ŏ	TYR	A 297	25.10	5 45.172	50.203		AAAA
MOTA	2361	N		A 298	24.77	2 44.989	52.417	1.00 22.47	AAAA
ATOM	2362	CA		A 298	24.57	2 46.415	52.566	1.00 24.27	AAAA
MOTA	2362	CB	HTS	A 298	23.46	R 46.726	53.556	1.00 23.17	AAAA
MOTA		CG	HTS	A 298	23.09	7 48.166	5 - 53.572	1.00 23.20	AAAA
ATOM	2364	_	בדג כ	A 298	23.58	B 49.201	54.287	1.00 24.25	AAAA
ATOM	2365		HIS	A 298	22.19		52.680		AAAA
MOTA	2366		DIE I	A 298	22.15		52.848	1.00 23.31	AAAA
ATOM	2367		HTC	A 298	22.98				AAAA
MOTA	2368		בבון ב	A 298	25.88	·			AAAA
MOTA	2369		UTC	A 298	26.28			1.00 24.47	AAAA
MOTA	2370	_	22C	A 299	26.56				AAAA
MOTA	2371		PRO	A 299	26.17				AAAA
MOTA	2372				27.84			1.00 27.31	AAAA
MOTA	2373		PRO	A 299	28.15			1.00 27.04	AAAA
MOTA	2374		PRO	A 299	26.74			1.00 27.57	AAAA
MOTA	2375	_	PRO	A 299	27.82				AAAA
ATOM	2376	C	PRO	A 299	21.02				•
			•					•	

MOTA	2377	0	PRO	Α	299	28	.755	48.82	6 54.	939	1.00	.28.04	AAAA
ATOM	2378	N	TYR			26	.769	49.79	4 54.	452	1.00	27.04	AAAA
ATOM	2379	CA	TYR				.629	50.47		740	1.00	27.59	AAAA
	2380	СВ	TYR				.425	51.43		700	1.00	30.57	AAAA
MOTA		-	TYR				.516	52.59		718		32.91	AAAA
MOTA	2381	CG					.181	52.46		491		33.45	AAAA
ATOM -	2382		TYR									33.91	
MOTA	2383		TYR				.160	53.48		538			
ATOM	2384	CD2	TYR	A	300		.837	53.80		969		34.19	AAAA
MOTA	2385	CE2	TYR	A	300		.809	54.83		018		34.64	
ATOM	2386	CZ	TYR	A	300	25	.468	54.65		807		34.56	
MOTA	2387	OH	TYR	Α	300	25	.389	55.63	0 51.	844		36.05	
ATOM	2388	С	TYR	Α	300	26	.454	49.53	8 56.	93.6	1.00	26.48	AAAA
ATOM	2389	0	TYR	Α	300	. 27	.073	49.72	6 57.	979 -	1.00	25.81	AAAA
ATOM	2390	N	ALA	Α	301	25	.581	48.54	7 56.	791	1.00	25.41	AAAA
ATOM	2391	CA	ALA				.328	47.60	6 57.	865	1.00	24.64	AAAA
	2392	CB	ALA				.164	46.73		511	1.00	25.32	-AAAA
MOTA	2393	C	ALA				.568	46.77		067		25.53	
MOTA	_		ALA				7.030	46.56		194		26.39	
MOTA	2394	0					1.108	46.30		950		25.83	
	2395	Ŋ	LEU					45.50		926		26.32	
MOTA	2396	CA	LEU				3.323					27.38	
ATOM	2397	CB	LEU				3.782	45.37		479			
MOTA	2398	CG	LEU				0.081	44.72		.024		28.18	
ATOM	2399		LEU).119	44.84		.502		29.32	
ATOM	2400	CD2	LEU				1.296	45.38		613		27.38	
MOTA	2401	С	LEU	А	302		3.398	46.18		.764		26.41	
ATOM	2402	0	LEU	А	302	29	874	45.64	-	.755		26.62	
MOTA	2403	N	ALA	Α	303	29	756	47.39		. 353		26.50	
ATOM	2404	CA	ALA	A	303	30	778	48.17	6 58	.022		25.92	
MOTA	2405	CB	ALA	Α	303	31	1.001	49.47	5 57	. 277	1.00	25.24	
MOTA	2406	C	ALA			30	0.490	48.46	4 59	.487	1.00	26.03	AAAA
ATOM	2407	ō	ALA.			_	1.325	48.17	5 60	.340	1.00	26.95	AAAA
	2408	N	ARG				3.322	49.02		.792	1.00	25.29	AAAA
MOTA	2409	CA	ARG				3.999	49.35		.179	1.00	23.46	AAAA
MOTA	2410	CB			304		7.641	50.05		.291	1.00	23.78	AAAA
MOTA		-CG	ARG				7.553	51.45		.629		24.59	
MOTA	2411				304		5.302	52.22	•	.091		25.85	
ATOM	2412	CD	ARG				5.067	51.46		.869		27.54	
MOTA	2413	NE			304		3.978	51.54		.637		28.36	
ATOM	2414	CZ					3.957	52.36		.695		26.48	
MOTA	2415		ARG					50.79		.358		28.45	
MOTA	2416		ARG				2.910			.053		23.18	
ATOM	2417	С	ARG				3.991	48.11				22.26	
MOTA	2418	0	ARG				9.591	48.09		.135		23.20	
MOTA	2419	N			305		3.330	47.07		.560			
ATOM	2420	CA			305		3.200	45.81		.292		22.33	
MOTA	2421	CB			305		7.319	44.86		.516		22.17	
MOTA	2422	С			305		9.516	45.13		.621		22.27	
ATOM	2423	0			305		9.763	44.75		.760		22.48	
MOTA	2424	N	TRP	Α	306	3 (0.366	44.96		.620		22.57	
ATOM	2425	CA	TRP	A	306	3:	1.634	44.30	7 61	.861		21.28	
ATOM	2426	CB	TRP	Α	306	32	2.279	43.88	5 60	.553		21.07	
ATOM	2427	CG	TRP	Α	306	3:	1.703	42.61	18 60	.004		20.75	
MOTA	2428	CD2	TRP	А	306	3.3	1.886	42.10	3 58	. 683	1.00	19.5	AAAA
	2429	CE2	TRP	A	306		1.352	40.79		.668	1.00	19.18	AAAA 8
MOTA	2430	CE3	TRP	Δ	306		2.456	42.61		.510		19.59	
MOTA			TRP				1.071	41.63	2 60	.713		20.5	
ATOM	2431		TRP				0.864	40.53		.922		19.74	
MOTA	2432	NET	TRP	A	300			39.99		.524		19.18	
MOTA	2433	CZ2	TRP	A	300		1.368		-	.367		18.98	-
MOTA	2434	CZ3	TRP	A	200		2.474	41.81				19.2	
MOTA	2435		TRP	A	306		1.933	40.51	-	.388			
MOTA	2436	С	TRP	A	306		2.571	45.15		.674		20.80	
MOTA	2437	0	TRP	A	306		3.459	44.63		.341		20.5	
ATOM	2438	N	THR	A	307		2.373	46.47		.614		20.1	
MOTA	2439	CA	THR	A	307	3:	3.175	47.39		.407		20.5	
ATOM	2440	CB	THR	A	307		2.861	48.88	31 63	.045		21.0	
ATOM	2441	OG1	THR	A	307		3.329	49.15		.718		21.2	
MOTA	2442	CG2	THR	Ā	307		3.523	49.83		.030	1.0	20.0	9 AAAA
AIUN	6446								-				

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		_		707	32.8	E3 4	7.135	64.893	1.00 2	20.88	AAAA
MOTA	2443	C	THR A	307	33.7	_			1.00 2		AAAA
MOTA	2444		THR A		31.5		6.851		1.00 2		AAAA
ATOM	2445		LEU A		31.1		6.543		1.00 2		AAAA
MOTA	2446		LEU A		29.6		6.340	66.644	1.00 2		AAAA
ATOM	2447		LEU A		28.8		7.656	66.674	1.00 2		AAAA
MOTA	2448		LEU A		27.3		7.473	66.411	1.00		AAAA
MOTA	2449		LEU /		29.1		8.283	68.045	1.00		AAAA
MOTA	2450	_		A 308			5.284	67.052	1.00		AAAA
MOTA	2451	C		80E A	31.8 32.2		5.186	68.215	1.00		AAAA
ATOM	2452	0		A 308	32.0		4.310	66.165	1.00		AAAA
MOTA	2453	Ŋ		A 309	32.6		3.069	66.544	1.00	23.12	AAAA
ATOM	2454	CA		A 309	32.5		2.016	65.413	1.00	22.33	AAAA .
MOTA	2455	CB		A 309			0.787	65.827	1.00	21.76	AAAA
ATOM .	2456			A 309	33.3 31.3		1.678	65.061	1.00		AAAA
MOTA	2457	CGI	ILE A	A 309	30.3		11.037	66.166	1.00		AAAA
MOTA	2458			A 309			13.377	66.790		24.52	AAAA
MOTA	2459	С		A 309	34.1 34.1		12.828	67.709	1.00	25.72	AAAA
MOTA	2460	0		A 309	34.		14.253	65.957		24.70	AAAA
ATOM	2461	N		A 310			14.570	66.099		24.20	AAAA
MOTA	2462	CA		A 310	36.9 36.9		45.417	64.944	1.00	23.29	AAAA
MOTA	2463	CB		A 310			45.712	65.123	1.00	23.17	AAAA
MOTA	2464	CG		A 310	38.		44.752	65.257	1.00	21.36	AAAA
MOTA	2465	CD2		A 310	39. 40.		45.472	65.490	1.00	20,62	AAAA
MOTA	2466	CE2		A 310	39.		43.354	65.202	1.00	20.01	AAAA
MOTA	2467	CE3		A 310	39.		46.938	65.273	1.00	22.82	AAAA
MOTA	2468	CD1	TRP	A 310	39.		46.803	65.497		22.30	AAAA
MOTA	2469	NE1	TRP	A 310			44.845	65.668		19.91	AAAA
MOTA	2470			A 310	40.		42.734	65.381		19.08	AAAA
MOTA	2471	CZ3		A 310	41.	_	43.477	65.610		19.40	AAAA
MOTA	2472			A 310			45.279	67.411		25.26	AAAA
MOTA	2473	C		A 310			44.945	68.109	1.00	24.71	AAAA
ATOM	2474	0		A 310			46.247.		1.00	26.76	AAAA
MOTA	2475	N		A 311			46.975	69.007	1.00	27.89	AAAA
MOTA	2476	CA		A 311 A 311			48.081	69.113	1.00	28.98	AAAA
MOTA	2477	CB	CYS	A 311			49.462	67.991	1.00	31.89	AAAA
ATOM	2478	SG		A 311			46.043	70.212	1.00	27.51	AAAA
ATOM	2479	C		A 311			46.127	71.135		26.90	AAAA
ATOM	2480	И О		A 312			45.169	70.187	1.00	27.33	AAAA
MOTA	2481	CA	CLIT	A 312		246	44.210	71.250		28.03	AAAA
MOTA	2482	CB	GLU	A 312		106	43.287	70.850	1.00	28.55	AAAA
MOTA	2483	CG		A 312		903	43.333	71.741		28.93	AAAA
MOTA	2484	CD		A 312		232	42.958	73.154		29.78	AAAA
ATOM	2485	OE1		A 312		954	41.957	73.345	1.00	30.81	AAAA
MOTA	2486 2487	OE2		A 312		754	43.653	74.071	1.00	30.79	AAAA
MOTA	2488	C		A 312		463	43.357	71 514	1.00	28.91	AAAA
MOTA	2489	ō		A 312		822	43.110	72 662	1.00	30.57	AAAA
MOTA	2490	Ŋ		A 313	36.	.081	42.889	70.436		29.04	AAAA
MOTA	2491	CA		A 313	37	.266	42.045	70.516	1.00	28.87	AAAA
MOTA	2492	CB	LEU	A 313	37	.524	41.373	69.157	1.00	29.39	AAAA
MOTA	2493	CG		A 313		.548	40.311	68.644		30.32	AAAA
ATOM	2494			A 313	36	.910	39.872	67.215	1.00	30.26	AAAA
ATOM	2495	CD	UEU C	A 313		.582	39.114	69.593	1.00	30.42	AAAA
ATOM	_	_	LEU	A 313		.474	42.888		1.00	27.75	AAAA
ATOM	2496	Ö	LEU	A 313		.215	42.553	71.808	1.00	27.34	AAAA
ATOM	2497 2498		SER	A 314		.642	43.986		1.00	27.95	AAAA
MOTA				A 314		.736	44.927	70.376		28.62	AAAA
MOTA	2499	CA CB		A 314		.690	45.937			27.49	AAAA
MOTA	2500			A 314		.703	46.904		1.0	30.12	AAAA
ATOM	2501		250	A 314		.666	45.653			29.67	AAAA
MOTA	2502		ದವಿ ಬಡು	A 314		.488	46.517		1.0	0 29.00	AAAA
ATOM	2503		رت.V	A 315		.676	45.302		1.0	0 30.78	AAAA
ATOM	2504		CT.V	A 315		.535	45.935			0 32.92	AAAA
MOTA	2505		تری ۲.۷	A 315		.542	47.452			0 34.92	AAAA
MOTA	2506		GI.V	A 315		.142	48.091		7 1.0	0 35.17	AAAA
ATOM	2507		250	A 316	37	.881	48.041			0 36.88	AAAA.
MOTA	. 2508	14	·					•			-
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ATOM	2509	CA	ARG	A	316	37.841	49.493	72.702	1.00	39.49	AAAA
ATOM	2510	CB	ARG	A	316	38.608	49.968	71.484	1.00	39.86	AAAA
MOTA	2511	CG	ARG	A	316	37.946	49.677	70.161	1.00	4077	AAAA
ATOM	2512	CD	ARG	A	316	38.843	50.226	69.077	1.00	41.47	AAAA
ATOM	2513	NE	ARG	Α	316	40.140	49.566	69.092	1.00	42.36	AAAA
MOTA	2514	CZ	ARG	Α	316	41.224	50.057	68.515	1.00	43.38	AAAA
ATOM	2515	NH1	ARG	А	316	41.159	51.217	67.882	1.00	44.76	AAAA
ATOM'	2516	NH2		•	316	42.361	49.385	68.556		43.71	AAAA
ATOM	2517	С			316	36.418	50.015	72.631		41.54	AAAA
MOTA	2518	ō			316	35.564	49.429	71.959		42.64	AAAA
ATOM	2519	N			317	36.163	51.119	73.329		43.10	AAAA
ATOM	2520	CA			317	34.830	51.720	73.356		44.51	AAAA
ATOM	2521	CB			317	34.809	52.936	74.293		46.17	AAAA
ATOM	2522	CG			317	34.472	52.614	75.759		49.65	AAAA
ATOM	2523	CD			317	35.426	51.623	76.439		52.51	AAAA
ATOM	2524		GLU			35.153	51.251	77.607		53.37	AAAA
ATOM	2525	OE2				36.444	51.214	75.831		54.14	AAAA
ATOM	2526	C			317	34.318	52.098	71.974		43.86	AAAA
ATOM	2527	ō	GLU			35.067	52.532	71.108		42.46	AAAA
ATOM	2528	N	VAL			33.023	51.916	71.779		44.79	AAAA
ATOM	2529	CA	VAL			32.394	52.197	70.502		45.57	AAAA
MOTA	2530	CB	VAL			31.098	51.369	70.324		45.36	AAAA
ATOM	2531		VAL			30.537	51.558	68.924		45.44	AAAA
MOTA	2532		VAL			31.366	49.911	70.612		46.35	AAAA
MOTA	2533	C	VAL			32.007	53.652	70.377		46.41	AAAA
ATOM	2534	Ō	VAL			31.199	54.145	71.165		46.53	AAAA
MOTA	2535	N	PRO	Α	319	32.584	54.370	69.396	1.00	46.89	AAAA
MOTA	2536	CD	PRO			33.581	54.017	68.375		46.44	AAAA
ATOM	2537	CA	PRO	A	319	32.209	55.774	69.247	1.00	47.62	AAAA
MOTA	2538	CB	PRO	A	319	33.022	56.206	68.024	1.00	46.96	AAAA
MOTA	2539	CG	PRO	Α	319	33.161	54.922	67.251	1.00	46.38	AAAA
MOTA	2540	С	PRO	A	319	30.709	55.743	68.977	1.00	48.64	AAAA
MOTA	2541	0	PRO	Α	319	30.236	54.860	68.262	1.00	48.61	AAAA
MOTA	2542	N	GLU	A	320	29.944	56667	69.544	1.00	49.24	AAAA
ATOM	2543	CA	GLU	А	320	28.522	56.598	69.288	1.00	50.01	AAAA
MOTA	2544	CB	GLU	Α	320	27.720	57.330	70.363	1.00	51.15	AAAA
MOTA	2545	CG	GĽU	А	320	27.828	58.831	70.339	1.00	53.01	AAAA
MOTA	2546	CD	GLU	A	320	26.825	59.474	71.282	1.00	54.34	AAAA
MOTA	2547	OE1	GLU	Α	320	25.604	59.273	71.077	1.00	54.04	AAAA
MOTA	2548	OE2	GLU	Α	320	27.255	60.171	72.228	1.00	55.06	AAAA
ATOM	2549	С	GLU	Α	320	28.206	57.168	67.921	1.00	49.78	AAAA
MOTA	2550	0	GLU	A	320	27.170	56.861	67.324	1.00	49.79	AAAA
MOTA	2551	N	LÝS	Α	321	29.116	57.980	67.407	1.00	49.26	AAAA
MOTA	2552	CA	LYS	Α	321	28.906	58.589	66.109		49.20	AAAA
MOTA	2553	CB	LYS	A	321	28.873	60.106	66.251	1.00	50.38	AAAA
MOTA	> 354	CG	LYS	A	321	30.234	60.674	66.634		52.88	AAAA
MÓTA	555	CD	LYS			30.717	60.180	68.002		53.76	AAAA
MOTA	2 <i>3</i> 56	CE	LYS	A	321	32.229	60.348	68.154		55.00	AAAA
MOTA	2557	NZ	LYS			32.715	61.725	67.829		55.95	AAAA
ATOM	2558	С	LYS			30.037	58.207	65.171		48.64	AAAA
MOTA	2559	Ο.	LYS			31.052	57.650	65.590		48.58	AAAA
MOTA	2560	N	LEU			29.854	58.511	63.894	1.00	47.78	AAAA
MOTA	2561	CA	LEU			30.870	58.238	62.896	1.00	46.13	AAAA
ATOM	2562	CB	LEU			30.248	57.638	61.638		46.84	AAAA
MOTA	2563	CG	LEU			29.240	56.504	61.848		47.71	AAAA
MOTA	2564		LEU			28.788	55.998	60.491	1.00	48.02	AAAA
MOTA	2565	CD2	LEU			29.853	55.374	62.667		48.21	AAAA
MOTA	2566	C	LEU			31.427	59.608	62.580		44.61	AAAA
MOTA	2567	0	LEU .			30.674	60.571	62.491		44.73	AAAA
MOTA	2568		ASN .			32.741	59.706	62.447		42.66	AAAA
MOTA	2569		ASN .			33.360	60.976	62.135	1.00	41.19	AAAA
MOTE	2570		ASN .			34.860	60.904	62.402		41.07	AAAA
MOTA	2571		ASN .			35.576	60.001	61.436		41.43	AAAA
ATOM ·	2572		ASN .			35.117	58.901	61.147		42.46	AAAA
MOTA	2573		ASN .			36.720	60.449	60.943		41.77	AAAA
ATOM	2574	С	ASN .	A	323	33.068	61.223	60.658	1.00	40.76	AAAA
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Figure 19-40

								•
	_			32.430	60.395	60.010	1.00 40.19	AAAA
MOTA	2575 0	ASN A 323				60.129	1.00 40.11	AAAA
ATOM	2576 N	ASN A 324		33.523		58.735	1.00 39.99	AAAA
MOTA	2577 CA	ASN A 324					1.00 39.54	AAAA
ATOM	2578 CB	ASN A 324					1.00 40.88	AAAA
ATOM	2579 CG	ASN A 324		33.003	65.114		1.00 40.77	AAAA
	2580 OD:	L ASN A 324		31.763	65.145	J J		AAAA
MOTA		2 ASN A 324		33.779	65.938		1.00 40.63	
ATOM		ASN A 324		33.918	61.786		1.00 40.10	AAAA
MOTA	2582 C	ASN A 324		33.320	61.468	56.678	1.00 39.24	AAAA
MOTA	2583 O	ASN A 324		35.144	61.376	58.011	1.00 40.41	AAAA
MOTA	2584 N	LYS A 325		35.908	60.519	57.126	1.00 41.41	AAAA
MOTA	2585 CA	LYS A 325	-		60.201	57.761	1.00 42.64	AAAA
MOTA	2586 CB	LYS A 325		37.262		56.828	1.00 44.45	AAAA
ATOM	2587 CG	LYS A 325		38.224	59.504	57.491	1.00 45.61	AAAA
ATOM	2588 CD	LYS A 325		39.575	59.199		1.00 45.88	AAAA
ATOM	2589 CE	LYS A 325		40.358	60.464	57.850	1.00 45.00	AAAA
	2590 NZ	LYS A 325		41.717	60.151	58.494		AAAA
ATOM	2591 C	LYS A 325		35.124	59.248	56.856	1.00 41.56	
MOTA		LYS A 325		35.042	58.781	55.716	1.00 41.35	AAAA
MOTA		ALA A 326		34.524	58.703	57.906	1.00 41.32	AAAA
MOTA	2593 N			33.732	57.492	57.774	1.00 41.07	AAAA
MOTA	2594 CA	ALA A 320		33.452	56.912	59.143	1.00 40.87	AAAA
MOTA	2595 CB			32.420	57.722	57.019	1.00 41.24	AAAA
MOTA	2596 C	ALA A 326			56.913	56.174	1.00 40.91	AAAA
ATOM	2597 0	ALA A 326		32.045		57.316	1.00 41.92	AAAA
ATOM	2598 N	LYS A 327		31.719	58.815	56.631	1.00 42.20	AAAA
ATOM	2599 CA	LYS A 327		30.451	59.097		1.00 43.61	AAAA
MOTA	2600 CE	LYS A 327		29.796	60.374	57.170	1.00 45.83	AAAA
	2601 CG			29.534	60.413	58,670	1.00 43.83	AAAA
MOTA	2602 CI			28.745	61.681	59.029	1.00 47.34	AAAA
MOTA				28.682	61.952	60.538	1.00 48.28	
MOTA		000		28.090	60.845	61.351	1.00 48.98	AAAA
MOTA		LYS A 327		30.673	59.266	55.125	1.00 41.33	AAAA
MOTA	2605 C	LYS A 327		29.879	58.797	54.309	1.00 40.78	AAAA
ATOM	2606 0	LYS A 327		31.761	59.950	54.781	1.00 40.39	AAAA
MOTA	2607 N	GLU A 328		32,129	60.217	53.399	1.00 38.91	AAAA
ATOM	2608 C				61.199	53.369	1.00 40.04	AAAA ·
ATOM	2609 CI	3 GLU A 328		33.300	62.576	53.909	1.00 41.94	AAAA
MOTA	2610 C	GLU A 328		32.941		53.994	1.00 43.77	AAAA
ATOM	2611 C	D GLU A 328		34.131	63.515	53.010	1.00 44.29	AAAA
ATOM	2612 0	E1 GLU A 328		34.904	63.595		1.00 45.11	AAAA
ATOM		E2 GLU A 328		34.285	64.189	55.040	1.00 37.39	AAAA
				32.497	58.938	52.675	1.00 37.33	AAAA
MOTA	2615 0			32.114	58.722	51.525	1.00 37.31	AAAA
MOTA	2616 N	2.2.2		33.255	58.091		1.00 35.67	AAAA
MOTA		200		33.657	56.820	52.783	1.00 33.03	
ATOM				34.451	56.012	53.813	1.00 30.62	AAAA
ATOM	2618 C			34.760	54.549		1.00 27.48	AAAA
MOTA	2619 C	G DEU A 325		35.549	54.453		1.00 36.24	AAAA
MOTA		D1 LEU A 329		35.514	53.936		1.00 25.74	AAAA
ATOM		D2 LEU A 329		32.405			1.00 33.24	AAAA
ATOM	2622 C	000		32.239			1.00 32.72	AAAA
ATOM	2623 O				_			AAAA
ATOM	2624 N			31.519			1.00 34.91	AAAA
MOTA		A LEU A 330		30.289				AAAA
ATOM		B LEU A 330		29.411				AAAA
MOTA	1627 C	G LEU A 330		30.067				AAAA
	. 2623 C	D1 LEU A 330		29.096	54.060			AAAA
MOTA	2629	D2 LEU A 330		30.512				AAAA
MOTA				29.499	55.695			
MOTA				28.984	54.968	3 51.060	1.00 36.14	AAAA
ATOM	2631 0			29.415			1.00 38.17	AAAA
MOTA				28.664			1.00 41.05	AAAA
ATOM		A LYS A 331		- 28.407			1.00 41.83	AAAA
MOTA	2634	B LYS A 331						AAAA
MOTA		G LYS A 331		27.584				AAAA
ATOM		D LYS A 331		27.202				AAAA
ATOM		E LYS A 331		26.182				AAAA
ATOM	-620	NZ LYS A 331		25.69				AAAA
	0000	LYS A 331		29.342		1 49.490		AAAA
ATOM		LYS A 331		28.71	2 57.98	0 48.480) T.OO 4T.34	, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
ATOM	2040							

	2641	NT	SER A 332	30.618	57.316	49.463	1.00 44.45	AAAA
MOTA	2641	N.		31.351	57.271	48.202	1.00 46.88	AAAA
MOTA	2642	CA	SER A 332			48.435	1.00 46.49	AAAA
MOTA	2643	CB	SER A 332	32.854	57.416			AAAA
MOTA	2644	OG	SER A 332	33.380	56.263	49.058	1.00 45.65	
MOTA	2645	С	SER A 332	31.093	55.959	47.494	1.00 48:73	AAAA
ATOM -	2646	0	SER A 332	31.262	55.854	46.281	1.00 49.51	AAAA
ATOM	2647	N	ILE A 333	30.697	54.952	48.258	1.00 50.62	. AAAA
ATOM	2648	CA	ILE A 333	30.420	53.648	47.686	1.00 52.65	AAAA
	2649	CB	ILE A.333	30.246	52.584	48.779	1.00 52.35	AAAA
MOŢA	2650		ILE A 333	29.889	51.248	48.157	1.00 51.40	AAAA
MOTA			ILE A 333	31.522	52.465	49.596	1.00 52.29	AAAA
MOTA	2651			31.403	51.463	50.696		AAAA
ATOM	2652		ILE A 333			46.924	1.00 54.42	AAAA
MOTA	2653	С	ILE A 333	- 29.120	53.712		1.00 55.10	AAAA
MOTA	2654	0	ILE A 333	28.122	54.178	47.462	1.00 56.56	AAAA
MOTA	2655	N	ASP A 334	29.118	53.274	45.672		
ATOM	2656	CA	ASP A 334	27.863	53.263	44.940	1.00 59.13	AAAA .
ATOM	2657	CB	ASP A 334	28.050	53.460	43.433	1.00 59.64	AAAA
ATOM	2658	CG	ASP A 334	28.976	52.446	42.823	1.00 59.23	AAAA
ATOM	2659	OD1	ASP A 334	28.853	52.194	41.606	1.00 58.87	AAAA
MOTA	2660		ASP A 334	29.839	51.925	43.559	1.00 59.34	AAAA
MOTA	2661	c	ASP A 334	27.251	51.898	45.215	1.00 60.95	AAAA
	2662	ō	ASP A 334	27.803	50.861	44.840	1.00 61.15	AAAA
MOTA			PHE A 335	26.113	51.914	45.897	1.00 62.56	AAAA
MOTA	2663	N		25.414	50.701	46.257	1.00 64.12	AAAA
MOTA	2664	CA	PHE A 335			47.779	1.00 64.40	AAAA
ATOM .	2665	CB	PHE A 335	25.311	50.621		1.00 64.98	AAAA
MOTA	2666	CG	PHE A 335	24.224	49.714	48.263	1.00 65.54	AAAA
MOTA	2667		PHE A 335	24.180	48.379	47.868		AAAA
MOTA	2668		PHE A 335	23.234	50.197	49.107	1.00 65.12	
ATOM	2669		PHE A 335	23.163	47.539	48.305	1.00 65.75	AAAA
MOTA	2670	CE2	PHE A 335	22.213	49.367	49.552	1.00 65.79	AAAA
ATOM	2671	CZ	PHE A 335	22.177	48.034	49.150	1.00 66.01	AAAA
ATOM	2672	Č.	PHE A 335	24.025	50.626	45.640	1.00 65.41	AAAA
ATOM	2673	ō	PHE A 335	23.591	49.564	45.184	1.00 65.27	AAAA
	2674	N	GLU A 336	23.338	51.763	45.618	1.00 66.38	AAAA
MOTA	_	CA	GLU A 336	21.980	51.826	45.097	1.00 67.49	AAAA
MOTA	2675		GLU A 336	21.893	51.260	43.673	1.00 68.25	AAAA
MOTA	2676	CB	GLU A 336	20.459	51.230	43.116	1.00 69.15	AAAA
MOTA	2677	CG		20.334	50.465	41.804	1.00 69.40	AAAA
ATOM	2678	CD	GLU A 336		49.271	41.784	1.00 69.57	AAAA
MOTA	2679		GLU A 336	20.710		40.804	1.00 69.10	AAAA
MOTA	2680		GLU A 336	19.851	51.051		1.00 67.68	AAAA
MOTA	2681	C	GLU A 336	21.098	50.999	46.025	1.00 67.58	AAAA
MOTA	2682	0	GLU A 336	21.216	49.776	46.082		
ATOM	2683	N	GLU A 337	20.227	51.679	46.761	1.00 67.87	AAAA
MOTA	2684	CA	GLU A 337	19.317	51.020	47.686	1.00 68.66	AAAA
ATOM	2685	CB	GLU A 337	18.583	52.085	48.502	1.00 68.88	AAAA
ATOM	2686	CG	GLU : 337	18.279	51.715	49.944		AAAA
MOTA	2687	CD	GLU A 337	19.527	51.587	50.789	1.00 67.70	AAAA
	2688		GLU 337	20.319	52.554	50.851	1.00 67.05	AAAA
ATOM	2689		GLU A 337	19.711	50.518	51.398	1.00 67.79	AAAA
MOTA		C	GLU A 337	18.322	50.222	46.827	1.00 69.28	AAAA
MOTA	2690		GLU A 337	17.886	50.705	45.780	1.00 69.50	AAAA
ATOM	2691	0		17.966	49.012	47.259	1.00 69.55	AAAA
MOTA	2692	N	PHE A 338		48.176	46.497	1.00 69.67	AAAA
MOTA	2693	CA	PHE A 338	17.035			1.00 70.51	AAAA
ATOM	2694	CB	PHE A 338	16.995	46.759	47.066		AAAA
MOTA	2695	CG	PHE A 338	16.225	45.789	46.221	1.00 71.57	
MOTA	2696	CD1	PHE A 338	16.666	45.462	44.936	1.00 72.04	AAAA
ATOM	2697	CD2	PHE A 338	15.052	45.208	46.698	1.00 71.69	AAAA
ATOM	2698		PHE A 338	15.944	44.566	44.138	1.00 72.23	AAAA
ATOM	2699		PHE A 338	14.323	44.313	45.909	1.00 71.93	AAAA
	2700	CZ	PHE A 338	14.770	43.991	44.627	1.00 72.11	AAAA
MOTA	2700	Č	PHE A 338	15.633	48.770	46.494	1.00 69.26	AAAA
MOTA			PHE A 338	15.072	49.029	45.434	1.00 68.86	AAAA
MOTA	2702	0	200 x 330	15.053	48.962	47.674	1.00 69.35	AAAA
MOTA	2703	N	ASP A 339		49.572	47.755	1.00 69.61	AAAA
MOTA	2704	CA	ASP A 339	13.733			1.00 69.48	AAAA
MOTA	2705	CB	ASP A 339	13.134	49.457	49.157	1.00 69.72	AAAA
MOTA	2706	CG	ASP A 339	11.819	50.233	49.299	1.00 03.72	- Anna

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- 500	2707	OT1 '	ASP A	339	11.813	51.462		1.00 69.39	AAAA
MOTA	2708	002	ASP A	339	10.790	49.618		1.00 69.78	AAAA
MOTA MOTA	2709		ASP A		13.972	51.035	47.440	1.00 69.95	AAAA
MOTA	2710	_	ASP A		14.305	51.815	48.333	1.00 69.92	AAAA AAAA
ATOM	2711		ASP A		13.810	51.389	46.168	1.00 70.23 1.00 70.39	AAAA
ATOM	2712		ASP A		14.023	52.748	45.699	1.00 70.59	AAAA
ATOM	2713		ASP A		12.757	53.283	45.041 43.791	1.00 70.86	AAAA
ATOM	2714		ASP A		12.397	52.517 51.302	43.791	1.00 70.93	AAAA
MOTA	2715	OD1	ASP A	340	12.126 12.399	53.125	42.699	1.00 70.89	AAAA
MOTA	2716		ASP A		14.482	53.674	46.807	1.00 70.63	AAAA
ATOM	2717	_	ASP F		15.688	53.847	47.008	1.00 71.13	AAAA
ATOM	2718 2719	N O	GLU A		13.543	54.259	47.544	1.00 69.95	AAAA ·
MOTA MOTA	2720	CA	GLU A		13.947	55.150	48.619	1.00 69.17	AAAA
MOTA	2721	CB		341	13.636	56.613	48.266	1.00 70.83	AAAA AAAA
MOTA	2722	CG	GLU 7	A 341	14.098	57.601	49.347	1.00 73.44 1.00 75.27	AAAA
MOTA	2723	CD		A 341	13.956	59.071	48.951 48.646	1.00 75.27	AAAA
ATOM	2724		GLU A		12.825	59.518 59.786	48.954	1.00 75.69	AAAA
ATOM	2725	OE2		A 341	14.984 13.367	54.819	49.983	1.00 67.09	AAAA
MOTA	2726	C		A 341 A 341	12.233	55.176	50.297	1.00 66.57	AAAA
MOTA	2727	0		A 342	14.158	54.114		1.00 64.87	AAAA
MOTA	2728 2729	N CA		A 342	13.767	53.779	52.148	1.00 62.55	AAAA
MOTA	2730	CB		A 342	14.265	52.377	52.589	1.00 62.81	AAAA
MOTA MOTA	2731	CG1	VAL	A 342	14.042	52.193	54.081	1.00 62.56	AAAA
ATOM	2732	CG2	VAL .	A 342	13.513	51.298	51.849	1.00 63.69 1.00 59.94	AAAA AAAA
ATOM	2733	С		A 342	14.483	54.822	52.982 54.054	1.00 59.91	AAAA
MOTA	2734	0		A 342	14.022	55.215 55.278	52.442	1.00 56.85	AAAA
MOTA	2735	N		A 343	15.609 16.457		53.085	1.00 54.01	AAAA
MOTA	2736	CA		A 343 A 343	15.639		53.605	1.00 54.18	AAAA
ATOM	2737 2738	CB CG		A 343	16.505		54.241	1.00 53.96	AAAA
MOTA	2739	OD1		A 343	15.947		54.785	1.00 54.59	AAAA
MOTA MOTA	2740		ASP	A 343 .	17.747		54.191	1.00 53.61 1.00 51.92	AAAA AAAA
ATOM	2741	С	ASP	A 343	17.186		54.242	1.00 51.92	AAAA
MOTA	2742	0		A 343	16.611		55.307 54.029	1.00 48.86	AAAA
MOTA	2743	N		A 344	18.458 19.240			1.00 45.59	AAAA
ATOM	2744	CA		A 344 A 344	19.847			1.00 43.94	AAAA
MOTA	2745 2746	CB CG		A 344	18.847			1.00 41.70	AAAA
MOTA	2747	CD		A 344	17.953			1.00 38.94	AAAA
MOTA MOTA	2748	NE		A 344	17.139			1.00 36.78	AAAA AAAA
ATOM	2749	CZ		A 344	16.176			1.00 34.81 1.00 34.11	AAAA
ATOM	2750	NH1		A 344	15.890			1.00 31.84	AAAA
MOTA	2751	NH2		A 344	15.506			1.00 44.83	AAAA
ATOM	2752	C		A 344	20.340 21.308			1.00 43.97	A AA
MOTA	2753	0		A 344 A 345	20.19			1.00 44.32	a: aa
ATOM	2754 2755	N CA		A 345	21.19	9 57.87 <i>7</i>	55.618	1.00 43.74	AAAA
MOTA	2756	CB		A 345	20.86	59.248	55.039	1.00 44.49	AAAA
ATOM ATOM	2757	OG		A 345	19.64			1.00 46.07	AAAA AAAA
ATOM	2758	C	SER	A 345	21.30		57.144	1.00 42.82 1.00 42.91	AAAA
ATOM	2759	0	SER	A 345	22.30				AAAA
MOTA	2760	N	TYR	A 346	20.28				AAAA
MOTA	2761	CA	TYR	A 346	20.29 18.94				. AAAA
ATOM	2762		TYR	A 346	18.63				AAAA
MOTA	2763	CG	TXK	A 346 A 346	19.29			1.00 38.74	AAAA
MOTA	2764	רבים נעט	LAND	A 346	19.02			1.00 37.71	AAAA
ATOM	2765 2766	(L)	AAA C	A 346	17.68	2 55.225	5 58.653	1.00 38.49	AAAA
ATOM	2767		TYR	A 346	17.40	5 53.882			AAAA
MOTA MOTA	2768		TYR	A 346	18.07	9 52.899		1.00 37.59	AAAA AAAA
MOTA	2769		TYR	A 346	17.79	_			AAAA
ATOM	2770) C		A 346	21.43				AAAA
MOTA	2771			A 346	21.96 21.80				AAAA
ATOM	2772	N	MET	A 347	21.80	. JJ.041			•
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MOTA	2773	CA	MET A	347	22.879	54.756	59.530	1.00 38.19	AAAA
MOTA	2774	CB	MET A		23.042	53.582	58.566	1.00 38.26	AAAA
MOTA	2775	CG	MET A		21.973	52.523	58,694	1.00 38.17	AAAA
ATOM	2776	SD	MET A		22.317	51.115	57.641	1.00 38.05	AAAA
		CE	MET A		22.237	51.892	56.101	1.00 37.61	AAAA
ATOM	2777				24.189	55.494	59.603	1.00 38.00	AAAA
ATOM	2778	C	MET A					1.00 37.40	AAAA
ayom	2779	0	MET A		25.127	55.033	60.250		
ATOM	2780	N	LEU A		24.248	56.637	58.929	1.00 38.08	AAAA
MOTA	2781	CA	LEU A		25.449	57.463	58.898	1.00 38.07	AAAA
MOTA	2782	CB	LEU A	348	25.445	58.330	57.638	1.00 36.66	AAAA
MOTA	2783	CG	LEU A	4 348	25.379	57.583	56.310	1.00 35.47	AAAA
MOTA	2784	CD1	LEU A	348	25.285	58.559	55.165	1.00 34.51	AAAA
ATOM	2785	CD2	LEU A	348	26.605	56.716	56.167	1.00 36.56 .	AAAA
MOTA	2786	С	LEU A	348	25.521	58.353	60.138	1.00 39.07	AAAA
ATOM	2787	ō	LEU A	348	26.546	58.980	60.406	1.00 38.81	AAAA
MOTA	2788	И.	GLU A		24.432	58.385	60.898	1.00 39.90	AAAA
	2789	CA	GLU 2		24.363	59,213	62.092	1.00 40.95	AAAA
ATOM		CB		A 349	22.961	59.821	62.203	1.00 41.70	AAAA
MOTA	2790				22.515	60.629	60.966	1.00 42.28	AAAA
MOTA	2791	CG		349			60.708	1.00 42.51	AAAA
ATOM ·	2792	CD		349	23.349	61.891		1.00 42.38	AAAA
ATOM	2793		GLU A		23.414	62.778	61.587		
MOTA	2794		GLU A		23.933	61.998	59.614	1.00 43.34	AAAA
ATOM	2795	С	GLU 2		24.740	58.511	63.406	1.00 41.12	AAAA
ATOM	2796	0	-	A 349	24.664	59.118	64.476	1.00 41.38	AAAA
MOTA	2797	N	THR A	350	25.140	57.243	63.326	1.00 40.86	AAAA
ATOM	2798	CA	THR 2	A 350	25.555	56.475	64.504	1.00 40.69	AAAA
MOTA	2799	CB	THR I	A 350	24.405	56.283	65.510	1.00 41.56	AAAA
ATOM	2800	CG1	THR A	A 350	24.062	57.54 <i>9</i>	66.078	1.00 41.48	AAAA
MOTA	2801	CG2	THR A	A 350	24.821	55.345	66.638	1.00 41.19	AAAA
ATOM	2802	С		A 350	26.109	55.109	64.141	1.00 40.14	AAAA
MOTA	2803	ō		A 350	25.857	54.595	63.055	1.00 39.93	AAAA
MOTA	2804	N		A 351	26.865	54.527	65.067	1.00 40.32	AAAA
ATOM	2805	CA		A 351	27.491	53.227	64.857	1.00 40.70	AAAA
	2806	CB		A 351	28.855	53.213	65.540	1.00 39.89	AAAA
ATOM	2807	CG	-	A 351	29.911	52.290	64.951	1.00 39.68	AAAA
ATOM			LEU		31.170	52.403	65.772	1.00 39.88	AAAA
MOTA	2808				29.414	50.861	64.945	1.00 40.70	AAAA
MOTA	2809		LEU A		26.612	52.091	65.384	1.00 41.12	AAAA
MOTA	2810	C		A 351	26.467	51.060	64.736	1.00 40.02	AAAA
MOTA	2811	0		A 351		52.292	66.567	1.00 42.99	AAAA
MOTA	2812	N		A 352	26.040			1.00 43.93	AAAA
MOTA	2813	CA		A 352	25.138	51.326	67.201		
ATCM	2814	CB		A 352	25.412	51.225	68.707	1.00 43.38	AAAA
MOTA	2815	CG		A 352	26.743	50.597	69.055	1.00 44.68	AAAA
MOTA	2816	CD		A 352	27.185	50.927	70.482	1.00 45.48	AAAA
MOTA	2817	CE		A 352	26.189	50.500	71.539	1.00 46.21	AAAA
ATOM	2818	NZ	LYS A	A 352	76.646	50.944	72.895	1.00 47.34	AAAA
MOTA	2819	С	LYS ?	A 352	23.723	51.838	67.003	1.00 44.40	AAAA
ATOM	2820	0	LYS A	A 352	23.375	52.917	67.488	1.00 45.79	AAAA
ATOM	2821	N	ASP A	A 353	22.904	51.083	66.287	1.00 44.78	AAAA
MOTA	2822	CA	ASP A	A 353	21.532	51.509	66.074	1.00 44.79	AAAA
ATOM	2823	CB		A 353	21.050	51.030	64.702	1.00 45.20	AAAA
MOTA	2824	CG	ASP .		21.146	49.544	64.546	.1.00 45.21	AAAA
	2825		ASP	353	21.806	49.086	63.581	1.00 45.06	AAAA
MOTA	2826		ASP		20.549	48.841	65.391	1.00 45.54	AAAA
MOTA			ACD :	A 353	20.645	50.993	67.217	1.00 44.44	AAAA
ATOM	2827	C	ASP A		21.042	50.113	67.973	1.00 44.29	AAAA
ATCM	2828	0				51.553	67.367	1.00 44.22	AAAA
ATOM	2829	N		A 354	19.439		66.550	1.00 44.38	AAAA
ATOM	2830	CD		A 354	18.839	52.617		1.00 44.18	AAAA
ATOM	2831	CA		A 354	18.500	51.163	68.419		
ATCM	2832	CB		A 354	17.371	52.170	68.238	1.00 44.52	AAAA
ATCM	2833	CG		A 354	17.368	52.341	66.749	1.00 44.66	AAAA
ATOM	2834	С		A 354	17.995	49.740		1.00 43.87	AAAA
ATOM	2835	0	PRO 2	A 354	17.962	49.152	67.249	1.00 44.48	AAAA
ATCM	2836	N	TRP .	A 355	17.588	49.198	69.469	1.00 43.23	AAAA
ATOM	2837	CA		A 355	17.051	47.851	69.500	1.00 42.88	AAAA
ATCM	2838	CB		A 355	16.743	47.401	70.927	1.00 46.42	AAAA.
A1 CH	2000						•		•

			x 25	: =	17.959	a 4	7.052	71.695	1.00	49.91	AAAA
ATOM	2839	CG :	TRP A 35		18.476		5.733	71.903	1.00	51.56	AAAA
MOTA	2840		TBP A 35		19.684		15.868	72.627		52.03	AAAA
ATOM	2841		TRP A 35		18.03	-	14.450	71.548		52.25	AAAA
MOTA	2842		TRP A 35 TRP A 35		18.84		7.915	72.284		50.53	AAAA
ATOM	2843		TRP A 35		19.88		7-, 208	72.846	1.00	51.63	AAAA
MOTA	2844	NE1	TRP A 3	55 E	20.46	_	14.763	73.003	1.00	52.64	AAAA
MOTA	2845		TRP A 3		18.81		13.352	71.921		53.12	AAAA
MOTA	2846		TRP A 3		20.00		43.518	72.642	1.00	53.02	AAAA
MOTA	2847		TRP A 3		15.78		47.767	68.675		40.28	AAAA
MOTA	2848	-	TRP A 3		15.01	-	48.720	68.591		39.82	AAAA
MOTA	2849		TRP A 3		15.59		46.610	68.065		36.83	AAAA
ATOM	2850		ARG A 3		14.44		46.365	67.225		33.70	AAAA
MOTA	2851	CA	ARG A 3	56	14.90	-	46.197	65.772		29.50	AAAA
MOTA	2852	CB	ARG A 3	20	15.63		47.423	65.256		25.22	AAAA
MOTA	2853		ARG A 3				47.194	63.973	1.00	21.53	AAAA
MOTA	2854		ARG A 3		16.41 17.05		48.435	63.533		18.55	AAAA
MOTA	2855		ARG A 3		17.97	_	48.533	62.574		17.06	AAAA
MOTA	2856	CZ	ARG A 3			-	47.451	61.919		17.64	AAAA
ATOM	2857	NHl	ARG A 3	56	18.40	-	49.721	62.241		11.56	AAAA
MOTA	2858		ARG A 3		18.44	-	45.095	67.773		34.63	AAAA
MOTA	2859	С	ARG A 3		13.83		44.117	67.051		35.86	AAAA
MOTA	2860	0	ARG A 3		13.60	_	45.112	69.079		34.58	AAAA
MOTA	2861	N	GLY A 3	57	13.58		43.112	69.734		34.33	AAAA
ATOM	2862	CA	GLY A 3	357	13.00		43.783	69.395		34.31	AAAA
MOTA	2863	С	GLY A 3		11.53		44.418	68.484		33.56	AAAA
ATOM	2864	0	GLY A 3		11.00		42.906	70.139		34.47	AAAA
MOTA	2865	N	GLY A 3	358	10.87		42.656	69.916		34.61	AAAA
MOTA	2866	CA	GLY A 3		9.46		41.389	70.655		34.47	AAAA
MOTA	2867	C	GLY A		996		40.821	71.345		34.27	AAAA
- MOTA	2868	0	GLY A 3		7.80		40.948	70.523		34.16	AAAA
MOTA	2869	N	GLU A		7.43		39.729	71.180		33.94	AAAA
MOTA	2870	CA	GLU A	359	5.9		39.644	71.174	1.00	34.78	AAAA
MOTA	2871	CB	GLU A	359	5.2		40.648	72.123		36.70	AAAA
MOTA	2872	CG	GLU A	359	3.8		41.020	71.740		38.40	AAAA
MOTA	2873	CD	GLU A	3 3 3 ,	3.0		40.108	71.600	1.00	39.65	AAAA
MOTA	2874	OE1	GLU A	333	3.5		42.234	71.584		38.52	AAAA
MOTA	2875		GLU A	339	8.0		38.549	70.464	1.00	32.86	AAAA
MOTA	2876	С	GLU A	309	8.6		38.692	69.427	1.00	32.92	AAAA
MOTA	2877	0	GLU A	355	7.9		37.375	71.036	1.0	0 32.63	AAAA
MOTA	2878	N	VAL A		8.4		36.215	70.409		0 32.70	AAAA
ATOM	2879	CA	VAL A		9.4		35.472	71.376	1.0	0 33.24	AAAA
MOTA	2880	CB	VAL A VAL A		10.0		34.252	70.701	1.0	0 32.99	AAAA
MOTA	2881				10.5	21	36.406	71.827	1.0	0 32.09	AAAA
MOTA	2882	CG2	VAL A VAL A		7.3		35.319	69.976		0 32.81	AAAA
ATOM	2883	C	VAL A		6.7		34.660	70.791		0 32.02	AAAA
MOTA	2884	0	ARG A		7.0		35.321	68.674		0 33.12	AAAA
MOTA	2885	N	ARG A		6.0	-	34.508	68.086		0 33.52	AAAA
MOTA	2886	CA	ARG A		6.1	_	34.558	66.565	1.0	0 33.43	AAAA
MOTA	2887	CB	ARG A		5.7		35.885		1.0	0 34.35	AAAA
MOTA	2888	CG	ARG A		6.0		35.972	64.469	1.0	0 33.90	AAAA
MOTA	2889	CD	ARG A	361	7.4		36.331		1.0	0 31.70	AAAA
ATOM	2890		ARG A	361	7.8		36.608		1.0	0 31.18	AAAA
ATOM	2891		ARG A	361	7.0		36.562	61.941	1.0	0 30.48	AAAA
ATOM	2892		ARG A	261	9.1		36.948		1.0	0 29.71	AAAA
MOTA	2893				6.0		33.057		1.0	0 34.20	AAAA
MOTA	2894	_	ARG A		7.1		32.537		1.0	0 33.79	AAAA
MOTA	2895		ARG A	367	4.9		32.407		1.0	0 34.68	AAAA
MOTA	2896		LYS A	362	4.8		31.022		1.0	0 35.62	AAAA
ATOM	2897		LYS A	363	3.3		30.555		1.0	08.78	AAAA
MOTA	2898		LYS A	362		378	31.226		1.0	0 40.38	AAAA
ATOM	2899		LYS A	362		505	32.777		1.0	0 42.09	AAAA
MOTA	2900		LYS A	362		208	33.446			00 41.94	AAAA
MOTA	2901		LYS A	362		473	34.909	68.45		00 39.85	AAAA
ATOM			LYS A	362		710	30.177		1.0	0 35.12	AAAA
ATCM		_	LYS A	362		425	29.30			00 34.14	AAAA
ATOM		. 0.	LYS A	302	0.	323	20.00.			- '	•
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			GLU		262	5.661	30.460	66.703	1.00 35.12	AAAA
MOTA	2905							65.699	1.00 35.62	AAAA
MOTA	2906	ÇA	GLU			6.445	29.741			
MOTA	2907	CB	GLU	Α	363	6.567	30.560	64.424	1.00 36.81	AAAA
ATOM	2908	CG	GLU			5.280	30.808	63.711	1.00 38.66	AAAA
			GLU			5.477	31.704	62.517	1.00 39.60	AAAA
ATOM	2909	CD						61.637	1.00 39.11	AAAA
ATOM -	2910		GLU			6.287	31.324			
ATOM	2911	OE2	GLU	Α	363	4.826	32.782	62.469	1.00 39.95	AAAA
	2912	C	GLU			7.836	29.450	66.181	1.00 35.14	AAAA
MOTA	-					8.321	28.316	66.098	1.00 34.50	AAAA
MOTA	2913	0	GLU							AAAA
ATOM	2914	N	LAY	Α	364	8.475	30.505	66.671	1.00 34.96	
ATOM	2915	CA	VAL	Α	364	9.830	30.431	67.180	1.00 34.44	`AAAA
			VAL			10.338	31.821	67.570	1.00 33.68	AAAA
ATOM	2916	CB						60 162	1.00 34.13	AAAA
MOTA	2917		VAL			11.739	31.722			
MOTA	2918	CG2	VAL	A	364	10.337	32.715	66.347	1.00 31.83	AAAA
	2919	С			364	9.908	29.499	68.370	1.00 34.44	AAAA
ATOM						10.789	28.640	68.430	1.00 36.01	-AAAA
MOTA	2920	0			364				1.00 33.27	AAAA
MOTA	2921	N	LYS	Α	365	8.980	29.649	69.305		
MOTA	2922	CA	LYS	A	365	8.970	28.790	70.476	1.00 33.20	AAAA
					365	7.968	29.319	71.508	1.00 34.28	AAAA
ATOM	2923	CB						72.033	1.00 33.67	AAAA
MOTA	2924	CG			365	8.307	30.705			
ATOM	2925	CD	LYS	Α	365	7.282	31.181	73.039	1.00 34.85	AAAA
	2926	CE	LYS	A	365	7.658	32.534	73.638	1.00 36.47	AAAA
MOTA						6.698		74.710	1.00 37.91	AAAA
ATOM	2927	NZ			365				1.00 32.75	AAAA
MOTA	2928	С	LYS	A	365	8.654		70.109		
ATOM	2929	0	LYS	A	365	9.071	26.421	70.818	1.00 31.95	AAAA
	2930	N			366	7.919	27.136	69.012	1.00 32.81	AAAA
MOTA						7.600		68.581	1.00 33.65	AAAA
MOTA	2931	CA			366				1.00 33.98	AAAA
MOTA	2932	CB	ASP	Α	366	6.459		67.557		
ATOM	2933	CG	ASP	Α	366	5.131	26.107	68.140	1.00 33.94	AAAA
			ASP			4.870	25.767	69.307	1.00 33.89	AAAA
MOTA	2934	ODI	ASF	•	300	4.332		67.412	1.00 35.08	AAAA
ATOM	2935	OD2	ASP						1.00 33.05	AAAA
ATOM	2936	C			366	8.820		67.940		
ATOM	2937	0	ASP	Α	366	9.140	24.006	68.172	1.00 33.66	AAAA
					367	9.473	25.959	67.102	1.00 33.07	AAAA
MOTA	2938	N							1.00 32.27	AAAA
MOTA	2939	CA			367	10.684				AAAA
ATOM	2940	CB	THR	Α	367	11.304	26.719		1.00 32.28	
ATOM	2941	OG1	THR	Α	367	10.473	27.039	64.520	1.00 30.64	AAAA
			THR			12.711		65.166	1.00 33.29	AAAA
MOTA	2942							67.442	1.00 31.71	AAAA
MOTA	2943	С	THR	A	367	11.680			1.00 30.45	AAAA
ATOM .	2944	0	THR	Α	367	12.178	23.918	67.352		
	2945	N	DEC.	Δ	368	11.955	25.896	68.426	1.00 32.05	AAAA
ATOM					368	12.888		69.482	1.00 32.49	AAAA
atom	2946	CA						70.421	1.00 32.27	AAAA
MOTA	2947	CB			368,					
ATOM	2948	CG	LEU	Α	368	14.097	27.809	69.960	1.00 32.71	
	2949	CD1	LEU	Α	368	15.488	27.170	69.899	1.00 33.00	AAAA
ATOM			LEU			13.709		68.597	1.00 31.86	AAAA
MOTA	2950								1.00 33.82	AAAA
ATOM	2951	С			368	12.455				AAAA
ATOM	2952	0	LEU	Α	368	13.266		70.489	1.00 34.29	
	2953	N	GLU	A	369	11.183	24.285	70.645	1.00 34.30	AAAA
ATOM					369	10.687		71.375	1.00 36.07	AAAA
MOTA	2954	CA						71.748	1.00 38.71	AAAA
ATOM	2955	CB	GLU	Α	369	9.211			1.00 38.71	
ATOM	2956	CG	GLU	A	369	8.974	24.285	72.920	1.00 40.18	AAAA
		CD			369	7.509	24.359	73.341	1.00 41.56	AAAA
MOTA	2957					6.917			1.00 41.83	AAAA
MOTA	.2958	OE1			369					AAAA
ATOM	2959	OE2	GLU	A	369	6.957			1.00 41.74	
MOTA	2960	С			369	10.893	21.822	70.611	1.00 37.16	AAAA
			CTI		360	11.338		71.196	1.00 37.00	AAAA
ATOM	2961	0	الراق	4	369				1.00 37.71	AAAA
ATOM	2962	N	LYS	Α	370	10.586				
	2963	CA			370	10.797	20.547	68.567	1.00 38.46	AAAA
ATOM					370	10.166		67.177	1.00 39.96	AAAA
MOTA	2964	СВ							1.00 42.68	AAAA
ATOM	2965	CG			370	8.646				AAAA
ATOM	2966	CD	LYS	A	370	8.092	20.320		1.00 44.54	
	2967	CE			370	6.572		65.781	1.00 45.55	AAAA
ATOM						6.009			1.00 45.50	AAAA
MOTA	2968	NZ	LYS	A	370				1.00 38.34	AAAA
ATOM	2969	С			370	12.282	20.235			
MOTA	2970	0			370	12.683	19.071	68.493	1.00 37.86	AAAA
ATOM	29,0	-		-		-		•		•

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					-					
- movie	2971	N i	ALA A	371	13	3.105	21.266		1.00 37.69	AAAA
	2971 2972		ALA A			1.543	21.057	68.226	1.00 37.20	AAAA AAAA
	2973		ALA A		15	5.258	22.375		1.00 35.48	AAAA
ATOM	2974	c i	ALA A	371		5.023	20.477	69.558	1.00 37.63 1.00 37.12	AAAA
ATOM	2975	0	ALA A	371		5.920	19.626	69.585	1.00 37.12	AAAA
ATOM	2976	N	LYS A	372		4.426	20.930	70.665 .71.995	1.00 37.46	AAAA
ATOM	2977		LYS A			4.796	20.432 21.156	73.095	1.00 36.52	AAAA
MOTA	2978		LYS A			4.022 4.287	22.634	73.111	1.00 22.67	AAAA
MOTA	2979		LYS A			3.309	23.396	74.022	1.00 22.67	AAAA
MOTA	2980	CE	LYS A	372		3.600	24.874	73.901	1.00 22.67	AAAA
MOTA	2981 2982		LYS A			2.692	25.708	74.785	1.00 22.67	AAAA
MOTA	2983		LYS A		1	4.495	18.957	72.077	1.00 37.60	AAAA AAAA
MOTA MOTA	2984		LYS A			5.367	18.171	72.407	1.00 37.26 1.00 38.17	AAAA
ATOM	2985	N	ALA A	373	_	3.249	18.595	71.789	1.00 39.55	AAAA
ATOM	2986	CA	ALA A			2.812	17.206	71.829 71.395	1.00 39.34	AAAA
ATOM	2987	CB	ALA A	A 373		1.365	17.109 16.277	70.972	1.00 41.02	AAAA
MOTA	2988	С	ALA A			3.675 4.366	15.410	71.561	1.00 42.26	AAAA
MOTA	2989	0	ALA A			3.663	16.416	69.725	1.00 42.06	AAAA
MOTA	2990		ALA Z	В 951		3.696	34.788	54.072	1.00 27.38	ZONE
HETATM	2991	ZN O1	SHA			4.578	33.295	53.458	1.00 31.95	SAHA
HETATM HETATM	2992	02	SHA	_		4.294	35:218	51.444	1.00 33.51	SAHA
HETATM	2994	N1	SHA	-	. 2	4.578	33.085	52.069	1.00 34.03 1.00 34.25	saha Saha
HETATM	2995	C1	SHA			4.063	34.053	51.246	1.00 34.23	SAHA
HETATM	2996	C2	SHA			23.090	33.625	50.259 48.816	1.00 39.33	SAHA
HETATM	2997	C3	SHA			23.548	33.781 33.274	47.852	1.00 40.86	SAHA
HETATM	2998	C4	SHA	_		22.498	34.413	47.455	1.00 43.37	SAHA
HETATM	2999	C5	SHA			21.061	34.017	46.092	1.00 46.72	SAHA
HETATM	3000	C6 C7	SHA SHA	-		19.754	34.714	45.787	1.00 48.75	SAHA
HETATM	3005	C8	SHA	-		19.960	35.720	44.693	1.00 50.75	SAHA
HETATM HETATM	1 3002	03	SHA		:	20.381	35.467	43.575	1.00 51.08	saha Saha
HETATM	3004	N2	SHA	_		19.591		45.085	1.00 52.52 1.00 54.25	SAHA
HETATM	3005	C9	SHA			19.842		44.507 45.215	1.00 55.76	SAHA
HETATM	1 3006		SHA			19.243		44.727	1.00 56.53	Saha
HETATM	1 3007		SHA			19.423 20.169			1.00 56.58	SAHA
HETATI	4 3008		SHA			20.165 20.755			1.00 55.93	SAHA
HETATI	4 3009	C14	SHA SHA			20.612			1.00 54.65	SAHA
HETATT	M 3010 M 3011		WAT			36.485			1.00 4.67	SOLV
HEIMI	4 3012		TAW			27.702				SOLV
HETATI	M 3013	OH2	TAW	D 4		23.251	30.387			SOLV
HETATI	M 3014		TAW			33.825				SOLV
HETATI	M 3015		TAW			24.866 34.145				SOLV
H. TAT	M 3016		TAW			7.921			1.00 20.79	SOLV
HAT	M 3017	OH2		_		17.863			1.00 28.94	SOLV
HE PAT	M 3018 M 3019	OH2				35.580	44.610			SOLV
HETAT	M 3020		TAW	D 11		49.208	3 27.797	65.303		SOLV
READY.	M 3020		TAW			20.490	34.049	61.067		SOLV
HETAT	м 3022	OH	TAW S	D 13		44.75		5 46.084 57.444		SOLV
HETAT	м 3023	OHZ	TAW S	D 14		22.457		65.163		SOLV
HETAT	м 3024	OH	TAW S	D 15		3.399			1.00 22.37	SOLV
HETAT	м 3025	OH	TAW S			32.273 26.329			1.00 27.86	SOLV
HETAT	M 3026		TAW S			48.24			3 1.00 15.09	SOLV
HETAT	M 3027		2 WAT			15.24		2 72.082		SOLV
HETAT	M 3028		2 WAT			26.44	4 9.269	9 52.63		SOLV
HETAT	M 3029		2 WAT	D 21		26.55	4 18.38			SOLV
HETAT	M 3031		2 WAT	D 22	•	39.45				SOLV
ואומה	M 3032		2 WAT	D 23		26.74				SOLV
HETAT	MM 3033	3 OH	2 WAT	D 24		44.66		8 39.06 3 70.66		
HETAT	rm 3034	i oh	2 WAT	D. 25		14.71		5 70.86 6 69.86		
HETAT	M 3035	он он	2 WAT	D 26		45.12 30.02		6 49.75		SOLV
HETA	rm 3036	5 OH	2 WAT	27 מי		30.02	- 1,.00	-		•

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HETATM	3037	OH2	WAT	D	28	20.659	28.788	43.520	1.00 28.55	SOLV
HETATM			WAT		29	32.271	38.000	53.512	1.00 47.72	SOLV
HETATM			WAT		30	18.285	29.333	54.536	1.00 21.34	SOLV
HETATM			TAW		31	49.978	38.669	73.461	1.00 31.02	SOLV
HETATM			WAT		32	21.587	50.386	71.043	1.00 14.52	SOLV
HETATM			WAT		33	46.784	32.121	33.375	1.00 31.79	SOLV
			WAT		34	33.359	39.755	49.117	1.00 16.13	SOLV
HETATM			WAT		35	7.687	37.657	51.568	1.00 27.22	SOLV
HETATM					36	44.238	35.392	33.961	1.00 19.67	SOLV
HETATM			WAT			10.908	25.384	58.206	1.00 33.51	SOLV
HETATM			TAW		37	36.758	27.243	70.552	1.00 39.61	SOLV
HETATM	3047		WAT		38		46.691	54.654	1.00 32.43	SOLV
HETATM	3048		TAW		39	45.825			1.00 32.43	SOLV
HETATM			WAT		40	52.489	20.282	52.165		SOLV
HETATM	3050		TAW		42	12.117	17.831	56.596	1.00 27.74	
HETATM			WAT		43	45.023	26.168	35.172	1.00 14.09	SOLV
HETATM			TAW		44	39.392	12.771	62.066	1.00 35.15	SOLV
HETATM	3053		TAW		45	3.930	26.970	63.814	1.00 22.23	SOLV
HETATM	3054		WAT		46	8.454	19.321	71.677	1.00 32.36	SOLV
HETATM	3055	OH2	$\mathbf{T}\mathbf{A}\mathbf{W}$	D	47	20.280	18.126	73.237	1.00 33.88	SOLV
HETATM	3056	OH2	TAW	D	48	9.321	39.409	54.873	1.00 18.57	SOLV
HETATM		OH2	WAT	D	49	50.852	41.323	58.048	1.00 21.25	SOLV
HETATM	3058	OH2	WAT	D	50	37.134	34.599	60.315	1.00 61.70	SOLV
HETATM			WAT		51	14.944	62.815	48.613	1.00 42.50	SOLV
HETATM			WAT		52	6.494	33.164	51.420	1.00 40.65	SOLV
HETATM			WAT		53	24.913	44.799	72.298	1.00 17.10	SOLV
HETATM			WAT		54	51.156	35.095	48.814	1.00 23.05	SOLV
HETATM			TAW		55	16.518	41.750	45.596	1.00 49.25	SOLV
HETATM			WAT		56	10.326	16.413	61.267	1.00 46.03	SOLV
HETATM			TAW		57	25.316	47.708	73.062	1.00 22.73	SOLV
HETATM			WAT		58	4.013	33.865	76.173	1.00 44.82	SOLV
			WAT		59	24.846	18.072	36.805	1.00 34.67	SOLV
HETATM			WAT		60	15.930	56.853	61.737	1.00 55.56	SOLV
HETATM			WAT		61	49.662	44.249	48.982	1.00 28.72	SOLV
HETATM	3009				62.	23.232	17.421	53.920	1.00 13.11	SOLV
HETATM			TAW			39.293	23.035	33.289	1.00 35.79	SOLV
HETATM			WAT		63	19.908	20.169	44.339	1.00 24.33	SOLV
HETATM			WAT		64	33.259	21.655	69.560	1.00 45.10	SOLV
HETATM			WAT		65	27.528	53.947	68.629	1.00 44.79	SOLV
HETATM			TAW		66		48.716	52.865	1.00 54.01	SOLV
HETATM			TAW		67	18.774		63.401	1.00 34.01	SOLV
HETATM			TAW		68	10.877	29.062	28.786	1.00 27.08	SOLV
HETATM		-	TAW		69	43.057	31.367		1.00 30.10	SOLV
HETATM			TAW		70	24.816	44.057	43.447		SOLV
HETATM			WAT		71	37.368	38.823	46.381	1.00 33.55	SOLV
HETATM			TAW		72	9.038	18.327	63.519	1.00 31.34	· SOLV
HETATM			TAW		73	51.799	20.829	65.265	1.00 28.32	
HETATM			TAW		74	17.556	58.515	57.254	1.00 19.27	SOLV
HETATM			WAT		75	28.436	27.904	79.425	1.00 27.13	SOLV
HETATM	3084		TAW		76	18.939	35.798	35.800	1.00 94.18	SOLV
HETATM	3085		WAT		77	34.359	31.251	46.688	1.00 73.70	SOLV
HETATM	3086		TAW		78	44.373	51.649	60.029	1.00 30.23	SOLV
HETATM			WAT		79	28.537	63.478	48.324	1.00 21.09	SOLV
HETATM	3088	OH2	WAT	D	80	6.869	44.113	72.030	1.00 28.59	SOLV
HETATM		OH2	WAT	ם	81	42.882	18.761	71.115	1.00 31.80	SOLV
HETATM	3090	OH2	WAT	D	82	36.712	59.078	53.901	1.00 40.11	SOLV
HETATM	3091	OH2	WAT	D	83	37.506	42.495	40.104	1.00 51.37	SOLV
HETATM	3092	OH2	WAT	D	84	40.054	38.439	55.415	1.00 20.07	SOLV
HETATM	3093		WAT		85	32.170	56.633	72.920	1.00 45.23	SOLV
HETATM	3094		WAT		86	24.470	53.877	47.119	1.00 41.18	SOLV
HETATM			WAT		87	48.585	35.663	67.518	1.00 33.40	. SOLV
HETATM	3006		WAT		88	29.541	57.166	42.788	1.00 44.61	SOLV
MIATAIM	3030		WAT		89	47.814	28.707	41.228	1.00 45.64	SOLV
HETATM	3000	0112	WAT	7	90	49.377	52.112	63.320	1.00 22.26	SOLV
HETATM	3000	0112	WAT	ה	91	44.219	43.589	43.912	1.00 39.90	SOLV
HETATM	3099	OHZ	WAT	ש	91	25.913	61.639	75.382	1.00 48.28	SOLV
HETATM	3100					8.623	30.749	49.707	1.00 40.37	SOLV
HETATM	3101		WAT		93				1.00 21.46	SOLV
HETATM	3102	OH2	WAT	ט	94	45.634	41.080	40.990	1.00 21.90	2020

	2102	OH2	tal N ITT	Ъ	95	29,984	34.886	51.725	1.00 35.75	SOLV
HETATM		OH2			96	13.051	21.934	49.804	1.00 46.73	SOLV
HETATM					97	32,412	65.913	55.822	1.00 43.39	SOLV
HETATM	3105		TAW		98	35.056	43.390	38.348	1.00 34.53	SOLV
HETATM	3106	OH2				22.360	47.680	60.688	1.00 19.16	SOLV
HETATM		OHZ	TAW	ע	99	50.755	19.722	57.906	1.00 42.45	SOLV
HETATM		OH2				7.875	37.690	74.094	1.00 37.18	SOLV
HETATM	3109	OH2	TAW	D	101		26.796	43.617	1.00 30.72	SOLV
HETATM	3110	OHZ	WAT	D	102	24.080	34.126	75.765	1.00 39.89	SOLV
HETATM	3111	OH2	WAT	D	103	45.206	54.786	40.685	1.00 29.58	SOLV
HETATM	3112	OH2	TAW	D	104	26.110		77.647	1.00 44.04	SOLV
HETATM	3113	OH2	WAT	D	105	25.918	39.658		1.00 42.22	SOLV
HETATM	3114	OH2	TAW	D	106	41.578	18.191	36.809 73.896	1.00 42.22	SOLV
HETATM	3115	OH2	TAW	D	107	31.945	51.420		1.00 41.13	SOLV
HETATM	3116	OH2	WAT	Ď	108	16.722	60.311	51.182		SOLV
HETATM	3117	OH2	WAT	D	109	43.604	38.573	78.141	1.00 36.22	SOLV
HETATM	3118	OH2	WAT	D	110	16.063	15.496	69.430	1.00 55.36	SOLV
HETATM	3119	OH2	WAT	D	111	21.630	22.785	49.145	1.00 36.52	
HETATM	3120	OH2	WAT	D	112	27.479	56.647	44.026	1.00 50.82	SOLV
HETATM	3121	OH2	WAT	D	113	14.739	51.674	61.674	1.00 35.55	SOLV
HETATM	3122	QH2	WAT	D	114	50.063	26.435	54.358	1.00 50.86	SOLV
HETATM	3123	OH2	WAT	D	115	43.935	38.427	73.129	1.00 44.21	SOLV
HETATM	3124	OH2	WAT	D	116	49.707	31.478	57.709	1.00 36.11	SOLV
HETATM	3125	OH2	WAT	D	117	25.032	43.463	55.676	1.00 38.06	SOLV
HETATM	3126	OH2	WAT	D	118	10.618	46.623	59.838	1.00 26.33	SOLV
HETATM	3127	OH2	WAT	D	119	48.466	33.382	61.437	1.00 19.82	SOLV
HETATM	3128	OH2	WAT	D	120	44.157	40.058	37.907	1.00 42.95	SOLV
HETATM	3129	OH2	WAT	D	121	51.267	29.446	52.889	1.00 38.93	SOLV
HETATM	3130	OH2	WAT	D	122	16.653	15.228	72.975	1.00 45.41	SOLV
HETATM	3131	OH2	WAT	D	123	36.898	45.148	41.936	1.00 27.00	SOLV
HETATM	3132	OH2	WAT	D	124	49.655	34.591	59.117	1.00 38.97	SOLV
HETATM	3133	OH2	TAW	D	125	12.285	57.594	42.107	1.00 23.56	SOLV
HETATM	3134	OH2	WAT	D	126	28.294	57.644	73.289	1.00 34.79	SOLV
HETATM	3135	OH2	WAT	D	127	19.138	60.403	61.551	1.00 28.58	SOLV
HETATM	3136	OH2	WAT	D	128	30.300	33.685	34.047	1.00 27.37	SOLV
HETATM	3137	OH2	WAT	D	129	40.898	53.983	47.254	1.00 16.30	SOLV
HETATM	3138	OH2	WAT	D	130	43.550	32.160	38.272	1.00 38.86	SOLV
HETATM	3139	OH2	WAT	D	131	18.624	13.959	56.194	1.00 37.70	SOLV.
HETATM	3140	OH2	WAT	D	132	18.580	12.901	62.894	1.00 27.28	SOLV
HETATM	3141	OH2	WAT	ם	133	35.830	30.296	50.621	1.00 42.47	SOLV
HETATM	3142	OH2	WAT	D	134	51.219	35.855	51.878	1.00 20.37	SOLV
HETATM	3143	OH2	WAT	ם י	135	50.428	22.486	49.267	1.00 39.37	SOLV
HETATM	3144	OH2	WAT	ם	136	51.633	29.369	63.918	1.00 33.99	SOLV
HETATM	3145	OH2	WAT	ם י	137	46.384	43.924	55.825		SOLV
HETATM		OH2	WAT	םי	138	30.356	25.767			SOLV
HETATM	3147	OH2	WAT	D	139	25.070	47.842			SOLV
HETATM	3149	2H2	WAT	Ď	140	47.097	49.394			SOLV
HETATM	3110	2H2	WAT	D	141	15.246	37.581			SOLV
HETATM	3150	2H2	WAT	ת מי	142	8.341	23.099	64.695		SOLV
HETATM	1.3121	SH2	WAT	ם י	143	30.065	18.220			SOLV
HETATM	1 2127	OH2	WAT	תי	144	11.930	46.453	57.606	1.00 36.15	SOLV
HETATM	3134	0112	*****	~						

INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/24700

A. CLASSIFICATION OF SUBJECT MATTER IPC(7) :C07K 14/00; G01N 35/573								
US CL :Please See Extra Sheet.								
According to International Patent Classification (IPC) or to both national classification and IPC								
B. FIELDS SEARCHED Minimum documentation searched (classification system follow	ed by classification symbols)							
U.S. : Please See Extra Sheet.	ed by classification symbolog	:						
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched								
Electronic data base consulted during the international search	(name of data base and, where practicabl	e, search terms used)						
Please See Extra Sheet.								
C. DOCUMENTS CONSIDERED TO BE RELEVANT								
Category* Citation of document, with indication, where a	ppropriate, of the relevant passages	Relevant to claim No.						
Y KAKUTA et al. Crystal Structure of the Human Heparan Sulfate N-Deacetyla Journal of Biological Chemistry. In Number 16, pages 10673-10676, see	se/N-Sulfotransferase 1. The 6 April 1999, Volume 274,	1-19						
Y SUEYOSHI et al. A role of Lys-614 of human heparan sulfate N-deacetul Letters. 1998, Volume 433, pages abstract.	1-19							
X Further documents are listed in the continuation of Box C. See patent family annex.								
Special categories of cited documents:	"T" later document published after the inte date and not in conflict with the app	emational filing date or priority lication but cited to understand						
"A" document defining the general state of the art which is not considered the principle or theory underlying the invention to be of particular relevance								
"E" earlier document published on or after the international filling date "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive sky when the document is taken alone								
cited to establish the publication date of another citation or other "Y" document of particular relevance; the claimed invention cannot be								
considered to involve an inventive step when the document is combined considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art								
"P" document published prior to the international filing date but later than the priority date claimed	family							
Date of the actual completion of the international search Date of mailing of the international search report								
29 DECEMBER 2000 25 JAN 2001								
Name and mailing address of the ISA/US Commissioner of Patents and Trademarks Box PCT Washington, D.C. 20231 Authorized officer ARDIN MARSOHEL								
Washington, D.C. 20231	ARDIN MARSHEL Telephone No. (703) 508-0196							

INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/24700

C-10	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Category*	Citation of document, with indication, where appropriate, of the relevant passages	
Y	AHMAD et al. WD Repeats of the p48 Subunit of Chicken Chromatin Assembly Factor-1 Required for in Vitro Interaction with Chicken Histone Deacetylase-2. The Journal of Biological Chemistry. 04 June 1999, Volume 274, Number 23, pages 16646- 16653, see especially the abstract	1-19
Y	JOHN et al. Rhizobium NodB protein involved in nodulation signal synthesis is a chitooligosaccharide deacetylase. Proceedings of the National Academy of Sciences, USA. January 1993, Volume 90, pages 625-629, see especially the abstract.	1-19
A.	US 5,780,594 A (CARTER) 14 July 1998, see the entire disclosure.	1-19
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INTERNATIONAL SEARCH REPORT

International application No. PCT/US00/24700

A.	CLASSIFICATION	OF	SUBJECT	MATTER
115	CT .			

550/550 and 455/7.2

B. FIELDS SEARCHED
Minimum documentation searched
Classification System: U.S.

530/300,333,350; 435/6,7.2; 514/9

B. FIELDS SEARCHED

Electronic data bases consulted (Name of data base and where practicable terms used):

CAS, BIOTECH ABS, MEDLINE, EMBASE, WPI, WEST covering search terms: deacetylase, human, crystal, histone, inhibitor, x-ray, and crystallography

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